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Prep & Review Time:

ACCESS DB # 165 465
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Scientific and Technical Information Center SEP 13 2005	•
SEARCH REQUEST FORM	
Requester's Full Name: TANE ZARA Examiner #: 77512 Date: 9-1	2-05
Art Unit: 1636 Phone Number: 2-0765 Serial Number: 09/896, 6	692
Location (Bldg/Room#): 2018 (Mailbox #): 2018 Results Format Preferred (circle): PAPER)	DISK
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To ensure an efficient and quality search, please attach a copy of the cover sheet, claims, and abstract or fill out the following	
Title of Invention: Novel HIV Specific OC,	<u>, 500.</u>
Inventors (please provide full names): S. AGAWAC	
Earliest Priority Date: 6/19/0/	
Search Topic: Please provide a detailed statement of the search topic, and describe as specifically as possible the subject matter to be searched, elected species or structures, keywords, synonyms, acronyms, and registry numbers, and combine with the concept or utility of the Define any terms that may have a special meaning. Give examples or relevant citations, authors, etc., if known.	Include the invention.
For Sequence Searches Only Please include all pertinent information (parent, child, divisional, or issued patent numbers) alona appropriate serial number.	ng with the
Please Search	
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Thanks.	
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STAFF USE ONLY Type of Search Vendors and cost where applicable	
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Searcher Location: Structure (#) Westlaw WWW/I	
Date Searcher Picked Up: 913/05Bibliographic	
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Maximum Match
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Perfect score:
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AZ976439 2M025JB24

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AZ579532 1M0367H10

AZ828544 2M0105004

A1439277 ti54402.x

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BH000100 2M0287A18

AZ807034 2M0069B05

BG927393 HNC1-1-F7

CD533465 31P16 Ara

AZ432757 1M0218L14

CF932153 SP1-AA (F6

CR753365 DKFZA69D

AZ371089 1M012ZD06

AZ652627 B012Z40w-

BQ587767 E012Z40w-

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1M0322C12	1M0231A01	E012442-0	DRABUA12	8015525-0	T. brucei	1M0502K14	1M0010L24		Arabidops	2M0110D12	2M0056P01	1M0016M18	1M0148A10	PRIO12a_B	1M0244L06	tz31g03.x	PRI016b_B	1M0146L22	PRI0147d	1M0435P20

ALIGNMENTS

	FEATURES BOUTCE	TITLE JOURNAL COMMENT	RESULT 1 AZB21337/c LOCUS DEFINITION ACCESSION VERSION VERSION KEYWORDS SOURCE ORGANISM REFERENCE
/organism="MMs musculus" /mol_type="genomic DNA" /mol_type="genomic DNA" /strain="C578L/6J" /db_xref="taxon:10090" /clone="UUGC2M0094I04" /sex="Male" /lab_host="B. Coli strain XL10-Gold, T1-resistant, F-" /clone lib="Mouse 10kb plasmid UUGCIM library" /note="Vector: PMD42nv, Purified genomic DNA from M. musculus C578L/6J (male) was obtained from the Jackson Laboratory Mouse DNA Resource (http://www.jax.org/resources/documents/dnares/). The DNA	Tel: 801 585 5606 Fax: 801 585 7177 Email: ddunn@genetics.utah.edu Insert Length: 10000 Std Error: 0.00 Plate: 0094 row: L column: 04 Seq primer: CGTTGTAAAACGACGGCCAGT Class: plasmid ends Location/Qualifiers 122	Junn, D., Adyagi, A., Barber, M., Beacorn, T., Dival, B., Hamil, C., Islam, H., Longacre, S., Mahmoud, M., Meenen, B., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R. Mouse whole genome scaffolding with paired end reads from 10kb plasmid inserts Unpublished (2000) Contact: Robert B. Weiss University of Utah University of Utah Contact: Robert B. Weiss University of Utah Contact: Robert B. Weiss Contact: Robert B. Weiss Contact: Robert B. Weiss Contact: Robert B. Weiss University of Utah Contact: Robert B. Weiss Contact:	AZ821337 AZ821337 AZ821337 AZ821337 AZ821337 AZ821337 AZ821337 AZ821337 AZ821337.1 GI:12991245 GSS. Mus musculus (house mouse) Mus musculus (house mouse) Mus musculus (house mouse) Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Rodentia; Sciurognathi; Muridae; Mus. 1 (bases 1 to 22)

ORIGIN

Query Match

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Length

ORIGIN

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SOURCE
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                                                                                                                                                                                                                                                                                                               Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: (
Plate: 0251 row: B column: 24
Seq primer: CACAGGAAACAGCTATGACC
Class: plasmid ends
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University of Utah Genome Center
University of Utah
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Mus musculus (house mouse)
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                                                                                                                                                                                                                                                                                                                                                                                                                     Tel: 801 585 5606
Fax: 801 585 7177
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Unpublished (2000)
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Location/Qualifiers
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                 /lab_host="E. coli strain XL10-Gold, T1-resistant, F-" /Clone lib="Mouse 10kb plasmid UUGC2M library" /note="Vector: PWD42nv, Purified genomic DNA from M. musculus C57BL/6J (female) was obtained from the Jackson Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Biomedical
(http://www.jax.org/resources/documents/dnares/). The DNA
                                                                                                                                                /db_xref="taxon:10090"
/clone="UUGC2M0251B24"
                                                                                                                                                                                       /mol_type="genomic DNA"
/strain="C57BL/6J"
                                                                                                                                                                                                             organism="Mus musculus"
/mol_type="genomic יומאת"
                                                                                                                            sex="Female"
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Pred. No. 6.9e+05;
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was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pwn42 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pW042 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent B. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."

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Best Local
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University of Utah Genome Center
University of Utah
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                                                                                                                                                                                                                                                                                               Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0085 row: O column: 05
Seq primer: CGTTGTAAAACGACGGCCAGT
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Unpublished (2000)
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                                                                                                                                                                                                                                                                                                                                                                               Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mouse whole genome scaffolding with paired end reads from 10kb
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/lab_host="E. Coli strain XL10-Gold, T1-resistant, F /clone lib="Mouse 10kb plasmid UUGCIM library" /note="Vector: PMD42nv, Purified genomic DNA from M. musculus C57BL/6U (male) was obtained from the Jack Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                          Biomedical
                                                                                                                                                         /mol_type="genomic DNA"
/strain="C57BL/6J"
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                                                                                                                      'db_xref="taxon:10090"
'clone="UUGC2M0085005"
                                                                                                    sex="Male"
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                    from the Jackson
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(http://www.jax.org/resources/documents/dnares/). The DNA

was hydrodynamically sheared by repeated passage through a

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REFERENCE
AUTHORS
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Н
                                                                                                                                                                                                                                                                                                                                                                                      to give a tight size distribution (
4 kb). The v + i method used for the library construction is described in detail in Smith, H. and Venter, J.C. (Making small insert libraries for whole genome shotgun sequencing projects. Genome Sequencing: A Practical Approach, eds. M. Vaudin and B. Barrell, Oxford University Press, 1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Submitted (10-DEC-2000) Trypanosoma brucei genome sequencing project, Sanger Centre, The Wellcome Trust Genome Campus, Hinxton, Cambridge CB10 1SA, E-mail: barrell@sanger.ac.uk and
                                                                                                                                                                                                                                                                                                               Email: nelsayed@tigr.org
Details of T. brucei sequencing at the Sanger Centre are available at http://www.sanger.ac.uk/Projects/T_brucei/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 nhl@Banger.ac.uk
Constructed at the Institute for Genomic Research (TIGR),
Rockville, MD. Genomic DNA isolated from a cloned population of
Trypanosoma brucei (TREU927/4 GUTat 10.1) was mechanically sheared
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hall,N., Bowman,S., Lennard,N.J., Doggett,J., Atkin,R., Chillingworth,C., Ormond,D., Harris,B., El-Sayed,N., Hou,L., Melville,S.E., Rajandream,M.A. and Barrell,B.G.
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AL483585.1 GI:11849086
GSS
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Direct Submission
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TCGCACCCATCTCTC 15
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                                              Conservative
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                                                                                                                                                                                /organism="Trypanosoma brucei"
|mol type="genomic DNA"
|strāin="TREU927"
|db_xref="taxon:5691"
                                                                                                                                                                                                                                                                                                  ocation/Qualifiers
                                                                  53.6%;
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                                                                  Score 11.8; DB 9;
Pred. No. 1e+06;
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                                                                                                                                                                                                                                                                13;
AZ579532 20 bp DNA linear GSS 13-DEC-20 1M0367H10F Mouse 10kb plasmid UUGC1M library Mus musculus genomic clone UUGC1M0367H10 F, genomic survey sequence.
                                                                                                                                                                                                                4 CACCCATCTCTCTCT 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Email: weisshaa@mpiz-koeln.mpg.de
Insert Length: 16 Std Error: 0.00
Plate: 15 row: N column: 03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Weisshaar B
ADIS DNA core facility at MPIZ
Max-Planck-Institute for Plant Breeding Research
Carl-von-Linne Weg 10, 50829 Koeln, Germany
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Herwig,R., Schulz,B., Weisshaar,B., Hennig,S., Steinfath,M., Drungowski,M., Stahl,D., Wruck,W., Menze,A., O'Brien,J., Lehrach,H and Radelof,U.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Beta vulgaris
Beta vulgaris
Butaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta,
Spermatophyta, Magnoliophyta, eudicotyledons, core eudicots;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BQ588621 16 bp mRNA linear EST 06-DEC-20
E012562-024-015-N03-SP6 MPIZ-ADIS-024-storage root Beta vulgaris
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Seq primer: SP6; CATACGATTTAGGTGACACTATAG Location/Qualifiers
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                                                                                                                                                                     CAACTTTCTCTCTCCCT 16
                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                              Sp6-Sall-CCACGCGTCCG-5prime-cDNA-polyA-CC-NotI-T7; Note: Sequencing granted in the context of the GABI-Beet project, local PI: Dr. Katharina Schneider, coordinator: Prof. Christian Jung; Sequence submission managed by RZPD/GABI-Primary database: http://gabi.rzpd.de"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /clone lib="MPIZ-ADIS-024-storage root"
/note="Vector: pCMVSPORT6; Site 1: SalI; Site 2: NotI;
/note="Vector: pCMVSPORT6; Site 1: SalI; Site 2: NotI;
CDNA library from sugar beet, library provided by KWS
Kleinwanzlebener Saatzucht AG Einbeck, Germany, contact:
b.schulz@kws.de; cloning sites SalI-NotI, primer sites and
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/mol_type="mRNA"
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                                                                                                                                                                                                                                                                                      Score 11.2; DB 5;
Pred. No. 1.8e+06;
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AZ828544 20 bp DNA linear GSS 20-FEB-20 2M0105004R Mouse 10kb plasmid UUGCIM library Mus musculus genomic clone UUGC2M0105004 R, genomic survey sequence.
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Niederhausern, A. and Wright, D., Weiss, R.

Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Seq primer: CGTTGTAAAACGACGGCCAGT Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erry
Plate: 0367 row: H column: 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tel: 801 585 5606
Fax: 801 585 7177
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AZ579532.1 GI:11693961
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/clone lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: PWD42nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackson
Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWD42 (gil4732114|gb)AF129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
adaptored vector DNA, and transformed into
chemically-competent E. coli XL10-Gold (Stratagene) cells
and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /mol_type="genomic DNA"
/strain="C57BL/6J"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /clone="UUGC1M0367H10"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         db_xref="taxon:10090"
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Pred. No.
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2.2e+06;
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                                                                                                                                                                                                                                                                                                                                    Length 20;
                                                       GSS 20-FEB-2001
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RESULT 8 AI439277

DEFINITION

AI439277 22 bp mRNA linear EST 09-M ti54a02.x1 NCI CGAP Lym12 Homo sapiens cDNA clone IMAGE:2134 similar to TR:\(\overline{Q}\)17089 \(\overline{Q}\)17089 \(\overline{C}\)DLAGEN ; contains element MER22

p mRNA linear EST 09-MAR-1999 sapiens cDNA clone IMAGE:2134250 3'

Focus

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                                                                                                                              Local Similarity
19
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                                                1 TCGCACCCATCTCTCTCT 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: 0.00
Plate: 0105 row: 0 column: 04
Seq primer: CACACAGGAAACAGCTATGACC
Class: plasmid ends
High quality sequence stop: 20.
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University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus. 1 (bases 1 to 20)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             plasmid inserts
Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mus musculus
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AZ828544.1 GI:12998452
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                                                                                                      Conservative
                                                                                                                                                                                                                      Laboratory Mouse DNA Resource
(http://www.jax.org/resources/documents/dnares/). The DNA
was hydrodynamically sheared by repeated passage through a
0.005 inch orifice at constant velocity. The sheared DNA
was blunt end-repaired with T4 DNA polymerase and T4
polynucleotide kinase. Adaptor oligonucleotides were
ligated to the blunt ends in high molar excess. The
adaptored DNA was purified and size-selected for a 9.5 to
10.5 kb range using preparative agarose gel
electrophoresis. Vector DNA was prepared from a derivative
of pWD42 (gi|4732114|gb|AF129072.1), a copy-number
inducible derivative of plasmid R1. The vector was ligated
with adaptors complementary to the insert adaptors and
purified. The sheared, adaptored mouse DNA was annealed to
adaptored vector DNA, and transformed into
chemically-competent E. coli XL10-Gold (Stratagene) cells
and selected for ampicillin resistance."
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/Clome_lib="Mouse 10kb plasmid UUGCIM library"
/note="Vector: PW942nv; Purified genomic DNA from M.
musculus C57BL/6J (male) was obtained from the Jackso
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /mol_type="genomic DNA"
/strain="C57BL/6J"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sex="Male"
                                                                                                                           50.0%;
73.7%;
                                                                                                   <u>,</u>
                                                                                                                              Score 11; DB 8;
Pred. No. 2.2e+06;
                                                                                                      Mismatches
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AI688330/c
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Best Local Similarity
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                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 22)

NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.

National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                AI688330 22 bp mRNA linear EST 17-DEC wc94c08.x1 NCI CGAP CO3 Homo sapiens cDNA clone IMAGE:2326286 similar to TR:Q64371 Q64371 PR-VBETA1. ;contains element MSR1
                                                    Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Elias Campo, M.D., Michael R.
                                                                                                                                                                                                                                                    Homo sapiens (human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA Sequencing by: Washington University Genome Sequencing Center Clone distribution: NCI-CGAP clone distribution information can be found through the I.M.A.G.B. Consortium/LLNL at: www-bio.llnl.gov/bbrp/image/image.html
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NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
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                                                                                                                  Tumor Gene Index
Unpublished (1997)
                                                                                                                                                                                                                                                                                                              AI688330.1 GI:4899624
                                                                                                                                                                                                                                                                                                                                                  repetitive element ;, mRNA sequence.
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Unpublished (1997)
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                                                                                            Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Life Technologies catalog #: 11547-015
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Robert Strausberg, Ph.D.
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cDNA Library Preparation: M. Bento Soares, Ph.D. cDNA Library Arraying: Greg Lennon, Ph.D.
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Location/Qualifiers
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                                          Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone_lib="NCI_CGAP_Lym12"
/note="Organ: lymph node, Vector: pCMV-SpORT6; Site_1:
Sall; Site_2: NotI; Cloned unidirectionally. Primer:
Oligo dT. Average insert size 1.25 kb. Life Technologies catalog #: 11547-015"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      tissue_type="lymphoma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            clone="IMAGE:2134250"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ab_host="DH10B"
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Pred. No.
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2.3e+06;
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                                                            Emmert-Buck,
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BH000100/c
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Email: ddunn@genetics.utah.edu Insert Length: 10000 Std Error: (Plate: 0287 row: A column: 18 Seq primer: CACACAGGAAACAGCTATGACC Contact: Robert B. University of Utah University of Utah Islam, H., Longacre, S., Mahmoud, M., Meenen, E., Pedersen, T., Reilly, M., Rose, M., Rose, R., Stokes, R., Tingey, A., von Niederhausern, A. and Wright, D., Weiss, R. BH000100 22 bp DNA linear GSS 27 2M0287A18R Mouse 10kb plasmid UUGC2M library Mus musculus clone UUGC2M0287A18 R, genomic survey sequence. Tel: 801 585 5606 Fax: 801 585 7177 Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Rodentia; SSD Mouse whole genome scaffolding with paired end reads from 10kb Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C., Mus musculus BH000100.1 84112, Rm. 308, Unpublished (2000) plasmid inserts Mus musculus (house mouse) .ass: plasmid ends (bases 1 to 22) quality sequence stop: 22.
Location/Qualifiers USA Biomedical GI:13871326 Weiss Genome Center Polymers Research Bldg., Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus. 0 00 20 S. 2030 E., 27-APR-2001 genomic SLC, S

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Best Local S
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source
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                                                                                                                                                                                                                                                                Contact: Robert B. University of Utah University of Utah
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19 bp DNA
2M0069B05R Mouse 10kb plasmid UUGCIM library Mus musculus genomic clone UUGCZM0069B05 R, genomic survey sequence.
                                                                                                    Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Erro
Plate: 0069 row: B column: 00
                                                                                                                                                                                                                                                                                                                                                                                                   Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T.,
Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von
Niederhausern,A. and Wright,D.,Weiss,R.
                                                                 Seq primer: CACACAGGAAACAGCTATGACC Class: plasmid ends
                                                                                                                                                                                                                                                                                                                                         Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Dunn, D., Aoyagi, A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mus musculus (house mouse)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AZ807034.1 GI:12970979
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:: 801 585 7177
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                                                                                                                                                                                                                                             308,
                     quality sequence stop: 19.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polymicleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent B. coll XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                           USA
                                                                                                                                                                                                                                           Biomedical Polymers Research Bldg., 20 S. 2030 E.,
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/clone_lib="Mouse_lokb plasmid_UUGC2M_library"
/note="Vector: PM912nv; Purified genomic_DNA from M.
musculus C57BL/6J (female) was obtained from the Jackson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (http://www.jax.org/resources/documents/dnares/). The DNA
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/mol_type="genomic DNA"
/strain="C57BL/6J"
/db_xref="taxon:10090"
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                                                                                                                                                                                                                                                                                            Genome Center
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                                                                                                                             Std Error: 0.00
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BG927393
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Matches

VERSION KEYWORDS

ACCESSION

Pocus

RESULT 11

SOURCE

FEATURES

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JOURNAL

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                                                                                                                                                                                                                                                                                                                                                              Identification and initial characterization of 5000 expressed sequenced tags (ESTs) each from adult human normal and osteoarthritic cartilage cDNA libraries Osteoarthr. Cartil. 9 (7), 641-653 (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 20)
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HNC1-1-F7.R HNC (Human Normal Cartilage) Homo sapiens cDNA, mRNA
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709 Swedeland Road,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sathe, G.
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                                                                                                                               Seq primer:
                                                                                                                                                                                   Fax: 610-270-5598
                                                                                                                                                                                                              Tel: 610-270-7245
                                                                                                                                                                                                                                                                                                             Contact: Sanjay Kumar
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                                                                                                                                       sanjay_kumar-1@gsk.com
imer: T7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ., Connor, J.R., Dodds, R.A., Halsey, W., Van Horn, M., Mui, P., Agarwal, P., Badger, A.M., Lee, J.C., Gowen,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /lab host="E. Coli strain XL10-Gold, T1-resistant, F-"/Clome_lib="Mouse 10kb plasmid UUGCIM library./note="Vector: PWPA2nv; Purified genomic DNA from M. musculus C57BL/6J (male) was obtained from the Jackso
/organism≈"Homo sapiens"
/mol_type="mRNA"
/db_xref="taxon:9606"
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/clone="UUGC2M0069B05"
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/strain="C57BL/6J"
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85.7%;
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                                                                                                                                                                                                                                 Box 1539, King of Prussia,
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RESULT 14
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Matches
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            119 Plant Science, Cornell University, Ithaca, NY 14853-5904, USA
Tel: 607 254 5418
Fax: 607 255 0599
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Susheng Gan
Department of Horticulture
Cornell University
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Arabidopsis thaliana
Arabidopsis thaliana
Bukaryota, Viridiplantae, Streptophyta; Embryophyta; Tracheophyta;
Eukaryota, Viridiplantae, Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Tosids; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
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Guo, Y., Cai, Z. and Gan, S.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: sg288@cornell.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CD533465.1 GI:40453477
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                                                                                                                                                          ATCTTTCTCTTTCT 16
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                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                        /tissue_type="Leaf"
/dev stage="Yellow Leaf With Greenish Base Area"
/lab_host="E. coli"
/lab_host="E. coli"
/lab_host="E. coli"
/lab_host="E. coli"
/lab_host=""E. coli"
/lab_host="E. co
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/lab host="E.coli DH10 B"
/clone lib="HNC (Human Normal Cartilage)"
/note="Vector: pSPORT I; Site_1: Sall; Site_2: NotI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /mol_type="mRNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           organism="Arabidopsis thaliana"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                db_xref="taxon:3702"
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                                                                                                                                                                                                                                                                                                           49.1%;
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Pred. No. 2.7e+06;
0; Mismatches 2;
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Pred. No. 2.7e+06;
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   GSS 03-OCT-2000
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Best Local Similarity
Matches 13; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AUTHORS
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AZ432757
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         High quality sequence stop: 19.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error:
Plate: 0218 row: L column: 14
Seq primer: CGTTGTAAAACGACGGCCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly,M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A. and Wright,D.,Weiss,R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Rodentia;
1 (bases 1 to 19)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          University of Utah Genome Center
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             plasmid inserts
Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mus musculus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AZ432757.1 GI:10556770
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CCTCTCCCTCTCCCTCT 3
                                            CCCATCTCTCTCTTCT 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ass: plasmid ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   308, Biomedical Polymers Research Bldg.,
                                                                                                    Conservative
                                                                                                                                                                                                                      (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWH2 (gi|4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    aboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note="Vector: PWD42nv; Purified genomic DNA from M.
nusculus C57BL/6J (male) was obtained from the Jack
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             lab_host="E. Coli strain XL10-Gold, T1-resistant, clone_lib="Mouse_10kb_plasmid_UUGCIM_library"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sex="Male"
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/strain="C57BL/6J"
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                                                                                                                        48.2%;
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                                                                                              Score 10.6; DB 8;
Pred. No. 3.3e+06;
0; Mismatches 4;
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                                                                                                                                                 Length 19;
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Search completed: September 15, 2005, 02:08:15
Job time : 1370.12 secs
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                                                                                                                                                                                                             Query Match 48.2%; Score 10.6; DB 7; Best Local Similarity 76.5%; Pred. No. 3.3e+06; Matches 13; Conservative 0; Mismatches 4;
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                                                                                                                                              2 CGCACCCATCTCTCTCC 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: marges@obstate.edu
Seq primer: REVERSE
High quality sequence stop: 20.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Margaret Essenberg
Biochemistry and Molecular Biology
Oklahoma State University
246 Noble Research Centre, Stillwater, OK 74078, USA
Tel: 4057446203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 (bases 1 to 20)
Patil,M.A., Pierce,M.L., Phillips,A.L., Venters,B. and
Essenby.K.
A pursuit of genes active in defense of upland cotton against
bacterial blight
Unpublished (2004)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gossypium hirsutum (upland cotton)
Gossypium hirsutum
Gossypium hirsutum
Gossypium hirsutum
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Tosids; eurosids II; Malvales; Malvaceae; Malvoideae; Gossypium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SP1-A4(P6-E4) A suppression subtractive hybridization cDNA library from bacterial-blight-resistant cotton Gossypium hirsutum cDNA clone SP1-A4(P6-E4), mRNA sequence. CF932153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CF932153.1 GI:45438734
EST.
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                                                                                                         CGCTCCCTTCCCTCCCC 17
                                                                                                                                                                                                                                                                                                                                                   /mol_type="mRNA"
/cultivar="Im216"
/db xref="taxon:3635"
/clone="SP1-A4(P6-E4)"
/clone="SP1-A4(P6-E4)"
/tissue_type="Foliar leaves inoculated with Xanthomonas
campestris_pv. malvacearum 3631"
/dev_stage="8, 14, 20, 30, 45 and 60 hours post
inoculation"
/clone_1lb="A suppression subtractive hybridization cDNA
library_from bacterial-blight-resistant cotton"
/note="Equal quantities of RNA from all stages were
combined_prior_to_mRNA isolation. In the cDNA subtraction
procedure, the reference cDNA was from non-inoculated
leaves"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           organism="Gossypium hirsutum"
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Minimum DB
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Listing first 45 summaries
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                                                                          a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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1: gb_ba:*

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4: gb_or

5: gb_or

6: gb_f

7: gb

9: gb

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            seq length: 0 seq length: 22
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Match Length DB
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1318.718 Million cell updates/sec
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ALIGNMENTS

VERSION KEYWORDS SOURCE ORGANISM REFERENCE	RESULT 2 BD077067 LOCUS DEFINITION ACCESSION	Matches Qy Db	ORIGIN Query Match Best Local	REFERENCE AUTHORS TITLE JOURNAL FEATURES SOURCE	RESULT 1 149132 LOCUS DEFINITION ACCESSION VERSION VERSION KEYWORDS SOURCE ORGANISM
BD077067.1 GI:22622670 JP 200151484-A/S. Human immunodeficiency virus Human immunodeficiency virus Viruses; Retroid viruses; Retroviridae; Lentivirus; Primate Lentivirus group. 1 (bases 1 to 22)	specific synthetic oligonucleotide and met	22; Conservative 0; Mismatches 0; Indels 1 TCGCACCCATCTCTCCTTCT 22 1	100.0%; Score 22; DB 6; Length 22; Similarity 100.0%; Pred. No. 4.6;	Unclassified. 1 (bases 1 to 22) 1 (bases 1 to 22) Cohen,A.S., Bourque,A. and Vilenchik,M. Method for analyzing oligonucleotide analogs Patent: US 5627277-A 6 06-MAY-1997; Patent: US 5627277-A 6 06-MAY-1997; 1. 1. 22 1. 22 /organism="unknown" /mol type="unassigned DNA"	149132 22 bp DNA linear F Sequence 6 from patent US 5627277. 149132 149132.1 GI:2467595 Unknown.
rimate	PAT 27-AUG-2002 hod of using	0; Gaps 0;			PAT 07-OCT-1997

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HYBRIDON INC
OS Human immunodeficiency virus
PN JP 2001514884-A/5
PD 18-SEP-2001
PF 05-AUG-1998 JP 2000509820
PR 19-AUG-1997 US 08/914827
PI SUDHIR AGRAWAL
PC C12N15/09,AG1K31/7125,AG1P31/18,C121
CCC Novel HIV-specific synthetic oligon
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1 (bases 1 to 21)
Cohen, A.S., Bourque, A. and Vilenchik, M. Method for analyzing oligonucleotide an Patent: US 5627277-A 7 06-MAY-1997;
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/mol_type="genomic DNA"
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           artificial sequences
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Agrawal, S.
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Potentiation of prodrug efficacy
Patent: WO 0134093-A 2 17-MAY-2001;
HYBRIDON, INC. (US)
Location/Qualifiers
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Human immunodeficiency virus
Viruses; Retroid viruses; Re
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nucleotides 1-4 and 18-21 and 2'0 methylribonucleosides"
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27-AUG-2002 of using

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HYBRIDON INC
OS Human immunodeficiency virus
PN JP 2001514884-A/2
PD 18-SEP-2001
PF 05-AUG-1998 JP 2000509820
PR 19-AUG-1997 US 08/914827
PI SUDHIR AGRAWAL
PC C12N15/09,A61K31/7125,A61P31/18,C12
CC Novel HIV-specific synthetic oligon
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BD077064.1 GI:22622667
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19-AUG-1997 US 08/914827
SUDHIR AGRAWAL
C12N15/09, A61K31/7125, A61P31/18, C12N15/00
Novel HIV-specific synthetic oligonucleotide
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18-SEP-2001
05-AUG-1998 JP 2000509820
19-AUG-1997 US 08/914827
SUDHIR AGRAWAL
C12N15/09,A61K31/7125,A61P31/18,C12N15/00
NOVEL HIV-specific synthetic oligonucleotide
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Location/Qualifiers
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/mol_type="genomic DNA"
/db_xref="taxon:12721"
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/mol_type="genomic DNA"
/db_xref="taxon:12721"
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BD077066
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            Patent JP 2001514884-A 4 18-SEP-2001;
HYBRIDON INC
OS Human immunodeficiency virus
PN JP 2001514884-A/4
PD 18-SEP-2001
PF 05-AUG-1998 JP 2000509820
PR 19-AUG-1997 US 08/914827
PI SUDDHIR AGRAWAL
PC C12N15/09,A61K31/7125,A61P31/18,C12N
CC Novel HIV-specific synthetic oligonu
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Human immunodeficiency virus
Viruses; Retroid viruses; Retroviridae;
lentivirus group.
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BD077065
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Novel HIV-specific synthetic oligonucleotide and method of using
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Novel HIV-specific synthetic oligonucleotide
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1 (bases 1 to 21)
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05-AUG-1998 JF 2000509820
19-AUG-1997 US 08/914827
SUDHIR AGRAWAL
C12N15/09,A61K31/7125,A61P31/18,C12N15/00
Novel HIV-specific synthetic oligonucleotide
using the same
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05-AUG-1998 JP 2000509820
19-AUG-1997 US 08/914827
SUDHIR AGRAWAL
C12N15/09,A61K31/7125,A61P31/18,C12N15/00
Novel HIV-specific synthetic oligonucleotide
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/mol_type="genomic DNA"
/db_xref="taxon:12721"
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RESULT 11
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OS Human im
PN JP 20015
PD 18-SEP-2
PF 05-AUG-1
PR 19-AUG-1
PR 19-AUG-1
PR C12N15/0
CC Novel HI
                                   Sequence
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Novel HIV-specific synthetic oligonucleotide
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JP 2001514884-A/6.
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Novel HIV-specific synthetic oligonucleotide
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19-AUG-1997 US 08/914827
SUDHIR AGRAWAL
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/mol_type="genomic DNA"
/db_xref="taxon:12721"
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/mol_type="genomic DNA"
/db_xref="taxon:12721"
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Building blocks with carbamate internucleoside
oligonucleotides derived therefrom
Patent: US 6605708-A 1 12-AUG-2003;
Location/Qualifiers
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Sequence 1 from patent US
AR374987
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Kandimalla, R. and Agrawal, S.
Cooperative Oligonucleotides
Patent: US 6372427-A 5 16-APR-2002;
Location/Qualifiers
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AR206325
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1 (bases 1 to 20)
Cohen, A.S., Bourque, A. and Vilenchik, M. Method for analyzing oligonucleotide an Patent: US 5627277-A 8 06-MAY-1997;
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1 (bases 1 to 21)
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AR170398
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Search completed: September 15, 2005, 01:23:27 Job time: 809.372 secs
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                                            Unclassified.
1 (bases 1 to 22)
Chang,L.-J.
Lentiviral vectors
Patent: US 6207455-A 8 27-MAR-2001;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                           Antiviral anticancer poly-substituted phenyl derivatized oligoribonuclectides and methods for their use Patent: US 6291438-A 24 18-SEP-2001; Location/Qualifiers
                                                                                                                                                                                                                                                             Unknown
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Sequence 8 from patent US 6207455.
AR140320
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Unclassified.
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Sequence 24 from patent US 6291438.
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AR170398.1 GI:17908357
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/mol_type="unassigned DNA"
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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AAT41969 AAT94860 AAZ88083 AAF85618	AAT41968 AAT41968 AAT94859 ADA18562 AAD22554	AAH23028 ACC59011 ADC24663 ABX90517	AAX19784 AAX19781 AAX19782 AAX19780 AAX19780	SUMMARIES ID
Aat41969 HIV-1 gag Aat94860 HIV-1 gag Aaz88083 Vector pT Aaf85618 HIV-1 vec		Aah23028 Control m Acc59011 Human Gag Adc24663 Antisense Abx90517 Mixed bac		Description Aax19779 Human imm Aax197783 Human imm

The present sequence represents a synthetic oligonucleotide sequence, antisense to a conserved gag region of the HIV-1 genome. The antisense oligonucleotide can be used to treat HIV-1 or HIV-2 infection in a

Claim 1; Page 64; 64pp; English.

New synthetic oligonucleotide sequences antisense to conserved gag region of ${\tt HIV-1}$ genome.

WPI; 1999-228890/19.

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Aad32574 Antisense	Aad32583 Antisense	Aad32581 Antisense	Aad32580 Antisense	Aad32573 Antisense	Aad32575 Antisense	Aad32578 Antisense	Aad32579 Antisense	Aad32585 Antisense	Aad32576 Antisense	Aax23632 Deletion	Aax18753 Target HI	Aav28224 Antisense	Aat10969 HIV gag c	Aaq61815 Antisense	Aaq43467 HIV viral	Adg36361 HIV siNA	Adg36335 HIV siNA	Adg35623 HIV sinA	Adg36315 HIV siNA	Adg36324 HIV siNA	Adg35577 HIV sina	Adg35597 HIV siNA	Adg35586 HIV sinA	The tadyon contains

ALIGNMENTS

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05-AUG-1998;
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                                                                                                                                                                                                             Synthetic.
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                                                                                                                                                                                                                                                Human immunodeficiency virus antisense oligonucleotide SEQ ID NO:5.
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                                                                             Agrawal S;
                                                                                                         19-AUG-1997;
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                                                                                                                                                    WO9909154-A2
                                                                                                                                                                                                      Human immunodeficiency virus 1.
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                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                       The present sequence represents a synthetic oligonucleotide sequence, antisense to a conserved gag region of the HIV-1 genome. The antisense oligonucleotide can be used to treat HIV-1 or HIV-2 infection in a mammal, or inhibit HIV-1 or HIV-2 in a cell. The oligonucleotide has less cell toxicity, provokes less immunostimulus than prior art, and is a GC-rich antisense HIV oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New synthetic oligonucleotide sequences antisense to conserved of {\tt HIV-1} genome.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BP; 2 A; 11 C; 1 G; 8 T; 0 U; 0 Other;
                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                97US-00914827
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/note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                 A; 11 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DNA;
                                                                                                                                  95.5%; but
100.0%; Pr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    100.0%;
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"phosphorothioate linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21
                                                                                                                                                                                                                                 1 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ₽P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
                                                                                                                                         Score 21; DB Pred. No. 17; O; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 22;
Pred. No.
                                                                                            22
                                                                                                                                                                                                                                 7 T; 0 U; 0 Other;
                                                                                                                                                                 DB 2;
17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
                                                                                                                                         0,
                                                                                                                                                                                  Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length
                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                         0,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         gag region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         less
GC-
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RESULT 4
AAX19701
IID AAX11
XX
AC AAX1
XX
DT 08-J
XX
DE Huma
XX
KW Huma
XX
KW infe
XX
OS Synt
OS Huma
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RESULT 3
AAX119784
ID AAX1
XX AAX1
AC AAX1
XX AAX1
XX AAX1
XX Huma
XX Huma
XX Huma
XX FH Key
FH Key
FH WO99
XX Synt
CO 25-F
XX W99
PM W999
XX Huma
XX W999
XX Huma
XX W999
XX C16-1
XX W91;
XX W91;
XX W91;
XX O16-1
XX W91;
XX O16-1
XX W91;
XX O16-1
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                                                                                                           Query Match
Best Local S
Matches 21
                                                                                                                                                                                                                                                                                                                                           The present sequence represents a synthetic oligonucleotide sequence, antisense to a conserved gag region of the HIV-1 genome. The antisense oligonucleotide can be used to treat HIV-1 or HIV-2 infection in a mammal, or inhibit HIV-1 or HIV-2 in a cell. The oligonucleotide has less
                                                                                                                                                                                                                            Sequence 21 BP; 2 A; 11 C; 1 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                               cell toxicity, provokes less immunostimulus than prior art, rich antisense HIV oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 28; Page 64; 64pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New synthetic oligonucleotide sequences antisense to conserved gag region of {\tt HIV-1} genome.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
Human immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human immunodeficiency virus antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAX19784;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1999-228890/19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Agrawal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   05-AUG-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-FEB-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WO9909154-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                08-JUN-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAX19784 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (HYBR-) HYBRIDON INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    nfection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    immunodeficiency virus; HIV; phosphorothioate linkage; gag;
tion; antisense oligonucleotide; ss.
                                                                                                                 21;
                                                                                                                                            Similarity
                             TCGCACCCATCTCTCTCCTTC 21
                                                                                                              ilarity 100.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             97US-00914827
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                                                                                                                                                                       95.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "phosphorothioate linkages"
                                                                                                                 0,
                                                                                                                                         Score 21; pred. No.
21
                                                                                                                 Mismatches
                                                                                                                                         DB
17;
                                                                                                                                                                    2;
                                                                                                                 0;
                                                                                                                                                                    Length
                                                                                                                 Indels
                                                                                                                                                                    21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQ ID NO:6.
                                                                                                                                                                                                                                                                                                                   and
                                                                                                              0;
                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                   is a
                                                                                                                                                                                                                                                                                                                   -59
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Human immunodeficiency virus; HIV; phosphorothioate linkage; gag; infection; antisense oligonucleotide; ss.

Human immunodeficiency virus 1.

Human

immunodeficiency virus

antisense oligonucleotide

SEQ ID

08-JUN-1999 AAX19781;

(first entry)

AAX19781 standard;

DNA;

21 ₽₽

0,

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AAX19782
ID AAX1
XX AAX1
AC AAX1
DT 08-J
XX Huma
XX Huma
XX Huma
XX Synt
OS Synt
OS Huma
XX Huma
XX Woos
FH Key
FT modi
FT modi
FT modi
FT woos
XX Woos
PN Woos
XX To-1
XX To-1
XX To-1
PF 05-1
XX To-1
PA (HYI
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present sequence represents a synthetic oligonucleotide sequence, antisense to a conserved gag region of the HIV-1 genome. The antisense oligonucleotide can be used to treat HIV-1 or HIV-2 infection in a mammal, or inhibit HIV-1 or HIV-2 in a cell. The oligonucleotide has less cell toxicity, provokes less immunostimulus than prior art, and is a GC-
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 21
                                                                                                                                                                                Synthetic
                                                                                                                                                                                                              Human immunodeficiency virus;
                                                                                                                                                                                                                                    Human
                                                                                                                                                                                                                                                          08-JUN-1999
                                                                                                                                                                                                                                                                                 AAX19782
                                                                                                                                                                                                                                                                                                      AAX19782 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Key
modified_base
                       19-AUG-1997;
                                                                                        WO9909154-A2
                                                                                                                                modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New synthetic oligonucleotide sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-AUG-1998;
                                                                  25-FEB-1999
                                                                                                                                                                    Human
                                                                                                                                                                                                  infection; antisense oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Agrawal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-FEB-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO9909154-A2
 (HYBR-) HYBRIDON INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (HYBR-) HYBRIDON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antisense
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1999-228890/19
                                                                                                                                                                   immunodeficiency virus
                                                                                                                                                                                                                                   immunodeficiency virus antisense oligonucleotide SEQ ID NO:2.
                                                                                                                                                                                                                                                                                                                                                                                   2 CGCACCCATCTCTCTCCTTCT 22
                                                                                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            genome.
                                                                                                                                                                                                                                                                                                                                                              CGCACCCATCTCTCTCCUUCU 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Page 63; 64pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 2 A; 11 C; 1 G; 4 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         provokes less immunostimulus than prior HIV oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               97US-00914827
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     98WO-US016345
                       97US-00914827
                                            98WO-US016345
                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                      DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                   95.5%;
85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "phosphorothioate linkages"
                                                                                                            "phosphorothioate linkages"
                                                                                                                                                                                                                                                                                                      21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      English.
                                                                                                                                                                                                                                                                                                      ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                   Score 21;
Pred. No.
                                                                                                                                                                                                               HIV;
                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                             phosphorothioate linkage; gag;
                                                                                                                                                                                                    88
                                                                                                                                                                                                                                                                                                                                                                                                                               DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        antisense
                                                                                                                                                                                                                                                                                                                                                                                                         0,
                                                                                                                                                                                                                                                                                                                                                                                                                              Length 21;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       to conserved
                                                                                                                                                                                                                                                                                                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       gag
                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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RESULT 6
AXX19780
ID AXX1
XX AXX1
XX AXX1
XX AXX1
XX Huma
XX H
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Best Local S
Matches 21
The present sequence represents a synthetic oligonuclectide sequence, antisense to a conserved gag region of the HIV-1 genome. The antisense oligonuclectide can be used to treat HIV-1 or HIV-2 infection in a mammal, or inhibit HIV-1 or HIV-2 in a cell. The oligonuclectide has less cannot be a conserved as a cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence represents a synthetic oligonucleotide sequence, antisense to a conserved gag region of the HIV-1 genome. The antisense oligonucleotide can be used to treat HIV-1 or HIV-2 infection in a mammal, or inhibit HIV-1 or HIV-2 in a cell. The oligonucleotide has less cell toxicity, provokes less immunostimulus than prior art, and is a GC-rich antisense HIV oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                   Claim 8;
                                                                                                                                                                                                                        New synthetic oligonucleotide sequences antisense to
                                                                                                                                                                                                                                                                         WPI; 1999-228890/19
                                                                                                                                                                                                                                                                                                                                                                                                                          19-AUG-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        05-AUG-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-FEB-1999.
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modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         08-JUN-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New synthetic oligonucleotide sequences antisense to conserved gag region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 1999-228890/19.
                                                                                                                                                                                                                                                                                                                        Agrawal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Agrawal
                                                                                                                                                                                                                                                                                                                                                                         (HYBR-) HYBRIDON INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          immunodeficiency virus; HIV; pho
tion; antisense oligonucleotide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     immunodeficiency virus antisense oligonucleotide SEQ ID NO:1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     immunodeficiency virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21;
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                                                                                                                                                                                                 genome.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21 BP; 2 A; 11 C; 1 G; 7 T; 0 U; 0 Other;
                                                                                                                                                   Page 63;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TCGCACCCATCTCTCTCCTTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first
                                                                                                                                                                                                                                                                                                                                                                                                                          97US-00914827
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           note=
                                                                                                                                                   64pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       95.5%; br
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "phosphorothioate linkages"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ø
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    phosphorothicate linkage; gag;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2
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                                                                                                                                                                                                                      conserved gag region
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RESULT 7
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Best Local S
Matches 18
UGACACCTGTTCTACUCAC-3'. Also described are: (1) potentiating the activity of a prodrug without producing side effects, comprising administering a polyanion before the prodrug; and (2) potentiating the activity of a prodrug without producing side effects, comprising co-administering a polyanion and a prodrug in a dosage which would not produce a therapeutic effect in the absence of the polyanion. The methods can be used for potentiating gents maximise the efficacy of the prodrugs, reducing the dosage to be administered to less toxic levels. The present sequence
                                                                                                         The present invention describes a method for potentiating prodrug activity, without producing side effects, comprising co-administering prodrug with a polyanion which is not an oligonucleotide having two 5' and four 3'2'-O-methylribonucleosides with the sequence: 5'-
                                                                                                                                                                                                Potentiating prodrug activity, without producing co-administering a prodrug with a polyanion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human immunodeficiency virus
                                                                                                                                                                         Example 1;
                                                                                                                                                                                                                                       WPI; 2001-300586/31
                                                                                                                                                                                                                                                                                                              09-NOV-1999;
                                                                                                                                                                                                                                                                                                                                      08-NOV-2000; 2000WO-US030687
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                                                                                                                                                                                                                                                                                                                                                                                                                                    misc_RNA
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13-AUG-2001
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                                                                                                                                                                                                                                                                                       (HYBR-) HYBRIDON INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 tumour;
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                                                                                                                                                                        Page 12; 26pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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(first en
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'note= "2'-0-methylribonucleosides'
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/note= "2'-O-methylribonucleosides"
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prodrug; polyanion; ss.
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Pred. No.
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RESULT 8
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Best Local &
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                                                                                                                                                                                                                                                     against VEGF for inhibiting cancer cell proliferation and angiogenesis. Preferably the oligonucleotide AAH23032 (a modified version of AAH22984) is used and may be utilized to treat Kapssi's sarcoma, ovarian cancer, prostrate cancer, pancreatic cancer or melanoma. The present sequence represents a control mixed backbone scrambled oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                       The invention provides a composition comprising one or more antisense Oligonucleotides directed against vascular endothelial growth factor (VEGF) where the antisense oligonucleotides inhibits proliferation of cells exhibiting autocrine VEGF activity at an IC 5 0 concentration of between 0.5-2.5 micro Ma. The antisense oligonucleotides may be directed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel antisense oligonucleotides useful for inhibiting vascular endothelial growth factor expression, angiogenesis and for treating cancer, e.g., Kaposi's sarcoma, ovarian cancer and prostrate cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   represents a phosphorothicate oligonucleotide complementary to HIV-1 gag which is used in an example from the present invention for the treatment of colon cancer tumour-bearing mice. (Updated on 11-SEP-2003 to
                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2001-451898/48.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gill PS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 13; Page 57; 105pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            antisense therapy;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Vascular endothelial growth factor; VEGF; antisense; angiogenesis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     proliferation; Kaposi's sarcoma; cancer; melanoma; cytostatic;
                                                                                                      18;
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                                                                                                      Conservative
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                                                                                                                                                                                                         G; 4 T; 3 U; 0 Other;
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Pred. No. 17;
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                                                                                                                                                     Length 21;
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                                                                                                Gaps
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RESULT 9 ACC59011

Human Gag/HIV-1 antisense oligonucleotide

01-JUL-2003

(first

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ACC59011;

ACC59011 standard; DNA;

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Best Local S
                                                                                                                                                                                                                             antiinflammatory, virucide, tuberculostatic, and protozoacide activity. The compounds may be useful in the manufacture of a medicament for the delivery of material that affects gene translation or gene transcription and modulates a biological process, in medical therapy. A compound is also useful for treating cancer, viral diseases such as infection caused by HIV, hepatitis (hepatitis B, hepatitis C and hepatitis D), herpes, TB, Epstein-Barr virus, malaria, influenza virus, Para influenza virus, molaria, influenza virus, para influenza virus, rhinoviruses, reoviruses, respiratory syncytial virus, rhinoviruses, polioviruses, coxsackie-viruses, echoviruses, rulalia virus, rhinoviruses, gastroenteritis viruses, riballa viruses, rulalia virus,
                                                                                                                                                                                                                                                                                                                                                                                                                    The invention relates to a novel compound comprising an optionally stabilised nucleac acid or its analogue encoding a peptide, protein or other biological modifier, aptamer, antisense sequence, or antisense mimic conjugated directly or through a linker to a ligand for the transcobalamin receptor or intrinsic factor receptor. A compound of the invention has cytostatic, antiviral, anti-HIV, hepatotropic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Compound useful for treating e.g. cancer comprises optionally stabilized nucleic acid, aptamer, antisense sequence, or antisense mimic conjugated to a ligand for the transcobalamin receptor or intrinsic factor receptor
                                                                                                                                             Sequence
                                                                                                                                                                                            molluseum vontagiosum virus, human parvovirus B19, cytomegalovirus, papillomavirus, varicella zoster, arenaviruses or filoviruses. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human; antisense; transcobalamin receptor; intrinsic factor receptor; cytostatic; antiviral; anti-HIV; hepatotropic; antiinflammatory; virucide; tuberculostatic; protozoacide; cancer; viral disease; ss; G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 88; 156pp; English.
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13-SEP-2002; 2002US-0410627P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (MAYO-) MAYO FOUND MEDICAL EDUCATION & RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2003-430085/40
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                                                                                       Similarity
                                                                                                                                             21
                                TCGCACCCATCTCTCTCTTC 21
                                                                        Conservative
                                                                                                                                       BP; 2 A; 11 C; 1 G; 4 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Callstrom
                                                                                                                                                                               used in the
                                                                                     95.5%;
85.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Prendergast FG;
                                                                                     Score 21;
Pred. No.
21
                                                                        Mismatches
                                                                                                                                                                           exemplification
                                                                                                        DB 8;
                                                                    0;
                                                                                                       Length 21;
                                                                                                                                                                             of.
                                                                      Indels
                                                                                                                                                                             the
                                                                                                                                                                             invention
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                                                                    Gaps
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RESULT 10 ADC24663

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07-NOV-2002

US2002165174-A1

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RESULT 11
ABX90517
ID ABX90
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                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                           The present invention relates to a cobalamin-bound detectable or therapeutic agent in combination with a cobalamin transport protein. In the manufacture of a medicament to increase the uptake of detectable agent useful in radioimaging or therapeutic agent for treatment of a disorder associated with abnormal cellular proliferation, an infectious disease and cardiovascular disorder; as an antibiotic or antiviral agent for transcription of all factor. The method increases efficiency of vitamin B12 or vitamin B12 conjugated materials. The presents sequence represents an antisense nucleotide that can be conjugated to the carrier
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Composition useful for the treatment of e.g. infectious disease comprises a cobalamin-bound detectable or therapeutic agent in combination with a cobalamin transport protein.
                                                                         Antisense; ss; angiogenesis; n
                                                                                                                Mixed backbone scrambled control antisense oligonucleotide
                                                                                                                                            01-MAY-2003
                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                       Antisense;
                                                                                                                                                                      ABX90517;
                                                                                                                                                                                               ABX90517 standard; DNA; 21
                                                                                                                                                                                                                                                                                                                                                                                                   described in the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 4; SEQ ID NO 11; 97pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-SEP-2001; 2001US-0326183P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO2003026674-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cardiovascular disorder; antibiotic; antiviral agent; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cobalamin-bound detectable; radioimaging; infectious disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antisense DNA #11 that can be conjugated to the carriers of invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18-DEC-2003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADC24663 standard; DNA; 21 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (MAYO-) MAYO
                                                                                                                                                                                                                                                                                                                       18;
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                                                                                                                                                                                                                                                                                                                                                                          21 BP; 2
                                                                                                                                                                                                                                                                                                                     Conservative
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                                                                                                                                          (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FOUND MEDICAL EDUCATION & RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (first entry
                                                                           probe; VEGF; vascular endothelial growth factor; neoplastic proliferation; cellular proliferation.
                                                                                                                                                                                                                                                                                                                                                                          A; 11 C; 1
                                                                                                                                                                                                                                                                                                                                95.5%;
                                                                                                                                                                                                 ВP
                                                                                                                                                                                                                                                                                                                                  Score 21;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                          G; 4 T;
                                                                                                                                                                                                                                                                                                                     Mismatches
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                                                                                                                                                                                                                                                                                                                                                                          0 Other;
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RESULT 12
AAT46110
ID AAT466
XX AAT46
AC AAT46
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Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to a composition comprising an antisense oligonucleotide directed against vascular endothelial growth factor (VEGF). The antisense oligonucleotide is useful for preparing a composition treating cancer, neoplastic proliferation, abnormal cellular proliferation and preventing angiogenesis. The present sequence is a
  Habus
                                                                                                                                                                                            09-FEB-1995.
                                                                                                                                                                                                                                                                                                                                modified_base
                                                                                                                                                                                                                                                                                                                                                                                               modified_base
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    carbamate; internucleoside; linkage; phosphodiester;
nuclease resistant; stable; stable duplex formation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Oligo 1 contg. carbamate internucleoside linkage.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              יייי ביישישועם comprising an antisense oligonucleotide directed against vascular endothelial growth factor, useful for preparing a composition for treating cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gill PS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30-JAN-1998; 98US-00016541.
19-JAN-2000; 2000US-00487023.
19-JAN-2001; 2001WO-US000019.
                                                                                               28-JUL-1993;
                                                                                                                                           27-JUL-1994;
                                                                                                                                                                                                                                        WO9504066-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25-FEB-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAT46110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAT46110 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 13; Page 20; 54pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (GILL/)
(MASO/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       31-JAN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      13-MAR-2001;
                                                (HYBR-) HYBRIDON INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               scrambled
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2003-255224/25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GILL P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TCGCACCCATCTCTCTCCTTC 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ucecacccarcrcrcrccouc 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Masood R;
Agrawal S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        control antisense oliginucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP; 2 A; 11 C; 1 G; 4 T; 3 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2001US-00805761.
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                                                                                               93US-00098945
                                                                                                                                             94WO-US008655
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/*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                    /*tag= b
/note= "3'-NH-COO-5'
                                                                                                                                                                                                                                                                                                                                                      note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA;
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                                                                                                                                                                                                                                                                                                                                                        "phosphodiester backbone"
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
                                                                                                                                                                                                                                                                                    (carbamate) linkage"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       phosphorothicate;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0,
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RESULT 13
AAT41968
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAT46110-16 are oligonucleotides (ON) contg. novel carbamate internucleoside linkages. Dinucleoside carbamates having the structure 5' nucleoside-3'-NHCOO-5'-nucleoside-3' are claimed as building blocks for ON synthesis. The carbamate internucleoside linkages provide increased resistance to nucleolytic degradation without hindering the ability of the ONs to undergo base stacking, and in prefd. configurations, without destabilising the double helix formed between the ONs and their
                                                                                                                                                                                                                                                                                                                                                           AAT41968
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 2; Page 18; 38pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         complementary target nucleic acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               carbamate to form
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1995-082177/11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              oligo:nucleotide building blocks -
bamate to form prods. with nuclease
                                                                                                                                                                                                                                                                                                                                                                                                                                        20;
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                                                                                                                                                                                                                                                                                                                                                                                                                      بر
                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                            standard; cDNA;
                                                                                                                                                                                                                                                                                                                                                                                                               TCGCACCCATCTCTCTCCTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21
                                                                                                                                                                                                                                                                                                                                                                                                     researciererer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP; 2 A; 11 C; 1 G; 7 T; 0
                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                              90.9%;
                                                                                                                                                                                                                                                                                                                                                            21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      English.
                                                                                                                                                                                                                                                                                                                                                            ВP
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 20;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                      21
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                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         u; o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               comprising a di:nucleoside resistance and stable duple
                                                                                                                                                                                                                                                                                                                                                                                                                                              DB
45;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length
                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               stable duplex
                                                                                                                                                                                                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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04-APR-1996;
                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                              gene function; ss
                                                                                                                                                                                                                                                                                                   Co-operative binding; duplex; antisense inhibition; target sequence; human immunodeficiency virus; HIV; dimerisation domain; T structure;
                                                                                                                                                                                                                                                                                                                                                                                                      24-JUN-1997
Kandimalla ER,
                                                                                12-APR-1995;
                                                                                                                                                            17-OCT-1996.
                                                                                                                                                                                                    WO9632474-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                           AAT41968;
                                         (HYBR-) HYBRIDON
                                                                                                                                                                                                                                                                                                                                                             gag binding oligonucleotide used as
                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
                                                                                95US-00420672
                                                                                                                     96WO-US004605
                                         INC
                                                                                                                                                                                                                                                                                                                                                             antisense
                                                                                                                                                                                                                                                                                                                                                               HIV inhibitor.
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AAT41965-T41994 are oligonucleotides (ON) that bind to HIV-1 mRNA and/or DNA and act as antisense inhibitors of HIV-1 gene expression. The ON are preferably used as duplexes, i.e. a first ON has a region that binds to an HIV-1 target sequence in a 5'-3' direction and a second region complementary to a second ON which has a first region which binds to the same target HIV-1 sequence but in a 3'-5' direction and a second region complementary to the first ON. Both ON bind to the HIV target sequence up

sequence up

Disclosure; Page 43; 84pp; English.

influenza virus.

Compsn. contg. at least two co-operative oligo:nucleotide(s) complementary to a target sequence - and with mutually compl dimerisation domains, for use as antisense inhibitors of HIV

mutually complementary hibitors of HIV and

WPI; 1996-477125/47.

Agrawal

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RESULT 14
AAT94859
ID AAT944859
XX AAT94
XX AAT94
XX AAT94
XX YAT94
XX YAT16
XX YAT17
XX YAT17
XX YAT17
XX YAT17
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XX AAT17
XX The 1
PT Olig(
CC COMP)
CC COMP
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Best Local S
Matches 20
The present oligonucleotide is an antisense oligonucleotide that binds to part of the gag gene of Human immunodeficiency virus type 1 (HIV-1) (ANT94853-54 and ANT94865). The oligonucleotide is used to exemplify the method of the invention. This method comprises two synthetic cooperative oligonucleotides, where each oligonucleotide comprises a region complementary to one of tandem, non-overlapping regions of a target nucleic acid, and a dimerisation domain at a terminus of each of the oligonucleotides. The dimerisation domains of the oligonucleotides are complementary to each other. The target nucleic acid is an mRNA, a single-stranded viral DNA, or a single-stranded viral RNA. The synthetic oligonucleotides can interact cooperatively to provide improved affinity, specificity, and biological activity as antisense molecules. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      gag gene;
synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   until a certain point along the target sequence, where the two binding ON are in close proximity and the remainder of the binding ON bind each other. ANT41965-741971 lack a dimerisation domain (domain hybridising to a second antisense ON) but these may be added to give extra stability to the inhibiting complex. The duplex/target site complex forms a T-shaped structure, inhibiting expression of HIV-1 nucleic acid. Also co-operative ON, when labelled, can be used to identify specific bacteria or viruses in cell cultures; to study function of specific genes as an alternative to use of 'knock out' animals. Co-operative ON have improved affinity and sequence specificity, reduced toxicity and better antisense activity compared with single, longer ON. Insertion of dimerisation domains into
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Anti:sense oligo:nucleotide compositions have at least 2 cooperative oligo:nucleotide(s) having a targetting and a dimerisation region - useful for inhibition of target nucleic acid expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HIV-1 gag
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAT94859 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1997-512714/47.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kandimalla ER,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           04-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               16-OCT-1997.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human immunodeficiency virus 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antisense
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    initiation codon region; target region; dimerisation domain; cooperative oligonucleotide; affinity; specificity; molecule; treatment; viral infection; influenza; HIV; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                Page 17; 91pp; English.
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Best Local
                                                                                      comprising a first and second synthetic cooperative oligonucleotides linked to their respective binding partner consisting of cyclodextrin, adamantane, streptavidin, or biottin. Each oligonucleotide has a region complementary to a tandem, non-overlapping region of a target nucleic acid that is separated by 0-3 bases. The oligonucleotide composition is useful as a pharmaceutical composition to inhibit the expression of nucleic acids in vitro, or for treating a mammal infected by HIV or by acquired immunodeficiency syndrome (AIDS). The cooperative oligonucleotides have improved sequence specificity for a single-stranded target, reduced toxicity, and improved biological activity as antisense molecules. The present sequence represents a cooperative oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               oligonucleotide composition used to, e.g. treat mammal infected by human immunodeficiency virus, comprises synthetic oligonucleotides linked to binding partner consisting of cyclodextrin, adamantane, streptavidin, or biotin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cooperative oligonucleotide; binding partner; cyclodextrin; adamantane; streptavidin; pharmaceutical composition; nucleic acids expression; HIV; acquired immunodeficiency syndrome; AIDS; anti-HIV; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kandimalla ER, Agrawal
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ALIGNMENTS

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DEFINITION
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Human immunodeficiency virus
Human immunodeficiency virus
Viruses; Retroid viruses; Retroviridae; Lentivirus; Primate
Lentivirus group.
1 (bases 1 to 21)
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BD077065.1 GI:22622668
JP 2001514884-A/3.
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Cohen,A.S., Bourque,A. and Vilenchik,M.
Cohen,A.S., Bourque,A. and Vilenchik,M.
Method for analyzing oligonucleotide analogs
Patent: US 5627277-A 7 06-MAY-1997;
Location/Qualifiers
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Sequence 7 from patent US
I49133
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Novel HIV-specific synthetic oligonucleotide and the same
Patent: JP 2001514884-A 3 18-SEP-2001;
HYBRIDON INC
OS Human immunodeficiency virus
P 2001514884-A/3
PD 18-SEP-2001
PF 05-AUG-1997 US 08/914827
PI SUDHIR AGRAWAL
PC C12N15/09,A61K31/7125,A61P31/18,C12N15/00
CC Novel HIV-specific synthetic oligonucleotic
                                                                                                                                                                               Patent: JP 2001514884-A 4 18-SEP-2001;
HYBRIDON INC
OS Human immunodeficiency virus
PN JP 2001514884-A/4
PD 18-SEP-2001
PF 05-AUG-1998 JP 2000509820
PR 19-AUG-1997 US 08/914827
PI SUDMIR AGRAWAL
PC C12N15/09, A61X31/7125, A61P31/18,C12N
CC Novel HIV-specific synthetic oligon
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1 (bases 1 to 21
Agrawal, S.
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JP 2001514884-A/4.
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Novel HIV-specific synthetic
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05-AUG-1998 JP 2000509820
19-AUG-1997 US 08/914827
SUDHIR AGRAWAL
C12N15/09,A61X31/7125,A61P31/18,C12N15/00
NOVEL HIV-specific synthetic oligonucleotide
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JP 2001514884-A/3
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05-AUG-1998 JP 2000509820
19-AUG-1997 US 08/914827
SUDHIR AGRAWAL
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/mol_type="genomic DNA"
/db_xref="taxon:12721"
                                                               /organism="Human immunodeficiency virus"
/mol_type="genomic DNA"
/db_xref="taxon:12721"
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OS Human im
PN JP 20015
PD 18-SEP-2
PD 18-SEP-2
PF 05-AUG-1
PR 19-AUG-1
PR SUDHIR A
PC C12N15/0
CC Novel HI
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I49132
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Human immunodeficiency virus
Viruses; Retroid viruses; Retroviridae;
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Cohen,A.S., Bourque,A. and Vilenchik,M.
Method for analyzing oligonucleotide an
Patent: US 5627277-A 6 06-MAY-1997;
                                                                                                                                                                                                                                                                         Patent: JP 2001514884-A 5 18-SEP-2001;
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JP 2001514884-A/5
18-SEP-2001
05-AUG-1998 JP 2000509820
19-AUG-1997 US 08/914827
SUDHIR AGRAWAL
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Novel HIV-specific synthetic oligonucleotide
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                                                                                       /organism='Human immunodeficiency virus'
                                     /organism="Human immunodeficiency virus'
/mol_type="genomic DNA"
/db_xref="taxon:12721"
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/mol_type="unassigned DNA"
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Patent: WO 0134093-A 2 17-MAY-2001;
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synthetic construct
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AX146648
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Cohen, A.S., Bourque, A. and Vilenchik, M.
Method for analyzing oligonucleotide analogs
Patent: US 5627277-A 8 06-MAY-1997;
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                                                                                                                                                                        /organism="synthetic construct"
/mol_type="unassigned DNA"
/db xref="taxon:32630"
/note="oligonucleotide sequence"
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from Patent WO0134093.
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Patent WO0152904.
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OS Human im
PN JP 20015
PD 18-SEP-2
PP 05-AUG-1
PR 19-AUG-1
PR SUDHIR A
PC C12N15/0
CC Novel HI
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Novel HIV-specific synthetic
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1 (bases 1 to 21)
Agrawal,S.
                                                                                                                                                                                                                                                                                                                                                                                                                         JP 2001514884-A/1.
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Location/Qualifiers
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AX203700.1
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                   CGCACCCATCTCTCTCCTTC 20
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05-AUG-1998 JP 2000509820
19-AUG-1997 US 08/914827
19-AUG-1997 US 08/914827
SUDHIR AGRAWAL
C12N15/09,A61K31/7125,A61P31/18,C12N15/00
C12N15/09,A61K31/7125,Tannon
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                                                                                                                                                                                                                                                                                          Human immunodeficiency virus
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/mol type="unassigned DNA"
/db_xref="taxon:32630"
/note="Scrambled mixed backbone oligonucleotides,
nucleotides 1-4 and 18-21 and 2'0 methylribonucleosides"
                                                                                                     /organism="Human immunodeficiency virus"
/mol_type="genomic DNA"
/db_xref="taxon:12721"
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Location/Qualifiers
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                                                            95.2%; Score 20; 100.0%; Pred. No.
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100.0%; Pred. No.
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Matches 20
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     Patent: JP 2001514884-A 6 18-SEP-2001;
HYBRIDON INC
OS Human immunodeficiency virus
PN JP 2001514884-A/6
PD 18-SEP-2001
PF 05-AUG-1998 JP 2000509820
PR 19-AUG-1997 US 08/914827
PI SUDHIR AGRAWAL
PC C12N15/09,A61K31/7125,A61P31/18,C12N
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HYBRIDON INC
OS Human immunodeficiency virus
PN JP 2001514884-A/2
PD 18-SEP-2001
PF 05-AUG-1998 JP 2000509820
PR 19-AUG-1997 US 08/914827
PI SUDHIR AGRAWAL
PC C12N15/09,A61K31/7125,A61P31/18,C121
CC Novel HIV-specific synthetic oligon
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Human immunodeficiency virus
Viruses; Retroid viruses; Retroviridae;
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Viruses; Retroid viruses; Retroviridae;
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Novel HIV-specific
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BD077068.1 GI:22622671
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1 (bases 1 to 21)
Agrawal,S.
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JP 2001514884-A/6

18-SEP-2001

05-AUG-1998 JP 2000509820

19-AUG-1997 US 08/914827

SUDHIR AGRAWAL

C12N15/09,A61K31/7125,A61P31/18,C12N15/00
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C12N15/09,A61K31/7125,A61P31/18,C12N15/00
Novel HIV-specific synthetic oligonucleotide
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/mol_type="genomic DNA"
/db_xref="taxon:12721"
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                                                                  1 (bases 1 to 21)

Kandimalla, B. R. and Agrawal, S.

Cooperative oligonucleotides

Patent: US 6372427-A 5 16-APR-2002;

Location/Qualifiers
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source
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/mol_type="genomic DNA"
/db_xref="taxon:12721"
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                                     organism="unknown"
|mol_type="unassigned
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DEFINITION
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AR374987
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AR140320/c
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Search completed: September 15, 2005, 01:23:26
Job time : 773.628 secs
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Matches 19
                                                                                    Query Match 87.6%; Score 18.4; DB 6; Length 22; Best Local Similarity 95.0%; Pred. No. 1.7e+02; Matches 19; Conservative 0; Mismatches 1; Indels
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                                                                                                                            Unknown.
Unknown.
Unclassified.
CE 1 (bases 1 to 22)

DRS (Chang,L.-J.
E Lentiviral vectors
1 to 227-MAR-2001;
Location/Qualifiers
1 . 22
"Tanism="unknown"
"unassigned DNA"
                                             20
                                                                                                                                                                                                                                                                          22 bp
Sequence 8 from patent US 6207455.
AR140320
AR140320.1 GI:14482816
                                                                                                                                                                                                                                                                                                                                                                                                                     ch 90.5%; Score 19; DB 6; 1 Similarity 100.0%; Pred. No. 82; 19; Conservative 0; Mismatches
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1 (bases 1 to 21)

Habus, I. and Agrawal, S.

Building blocks with carbamate internucleoside linkages and oligonucleotides derived therefrom Patent: US 6605708-A 1 12-AUG-2003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19;
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Sequence 1 from patent US 6605708.
AR374987
AR374987.1 GI:40078169
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                                                          CGCACCCATCTCTCTCCTTC 20
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/mol_type="mRNA"
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Maximum Match 100%
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Perfect score:
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      Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                          Score
    seq length: 0
seq length: 22
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Match
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1: /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
2: /cgn2_6/ptodata/1/ina/5B_COMB.seq:*
3: /cgn2_6/ptodata/1/ina/6A_COMB.seq:*
4: /cgn2_6/ptodata/1/ina/6B_COMB.seq:*
5: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
6: /cgn2_6/ptodata/1/ina/backfiles1.seq:*
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PCT-US95-00163-6
US-08-178-660-7
US-08-178-660-8
PCT-US95-00163-7
US-08-178-660-8
PCT-US95-00163-8
US-08-420-672-5
US-08-935-312-8
US-08-420-672-6
US-08-318-138-8
US-08-318-138-8
US-08-644-786-1
US-08-664-786-1
US-09-587-934B-19
US-09-587-934B-19
US-09-587-934B-19
US-08-42-664-32
US-08-484-138-32
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Query Match Query Match Query Matches 100.0%; Score 22; DB 1; Best Local Similarity 100.0%; Pred. No. 0.74; Matches 22; Conservative 0; Mismatches 0 Matches 17CGCACCCATCTCTCTCTCT 22	US-08-178-660-6 Sequence 6, Application US/08178660 Patent NO. 5627277 GENERAL INFORMATION: APPLICANT: Coben, Aharon S. APPLICANT: Bourque, Andre APPLICANT: Wilenchik, Maria TITLE OF INVENTION: Oligonucleotide Analogs INMBER OF SEQUENCES: 8 CORRESPONDENCE ADDRESS: ADDRESSEE: Lappin & Kusmer STREET: 200 State Street CITY: Boston STATE: Massachusetts COUNTRY: U.S.A. ZIP: 02109 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0, Version #1. CURRENT APPLICATION: 536 APTICATION: 536 APTICATION: 536 ATTORNEY/AGENT INFORMATION: NAME: Kerner, Ann-Louise REGISTRATION NUMBER: US/08/178,660 FILING DATE: REGISTRATION NUMBER: 33,523 REFERENCE/DOCKET NUMBER: HYZ-012 TELEPONNE: 617/330-1311 INFORMATION FOR SEQ ID NO: 6: SEQUENCE CHARACTERISTICS: LENGTH: 22 base pairs TYPE: mucleic acid STRANDENNESS: single TOPOLOGY: linear MCLECULE TYPE: CDNA US-08-178-660-6	C 28 17 77.3 20 3 US-09-074-357-9 29 17 77.3 20 3 US-09-371-774-50 30 17 77.3 20 5 PCT-US95-06161-2 31 17 77.3 20 5 PCT-US95-06161-3 32 16.8 76.4 21 3 US-08-473-096-22 34 15.4 70.0 20 3 US-09-166-186-51 35 15.4 70.0 20 3 US-09-166-186-51 20 36 15 68.2 18 1 US-08-473-096-20 21 39 14.4 65.5 19 1 US-08-473-096-20 38 14.4 65.5 19 1 US-08-473-096-20 39 14.2 64.5 19 3 US-08-0871-579-1 40 14.2 64.5 21 3 US-08-971-579-1 41 14.2 64.5 21 1 US-08-981-194A-24 42 14.2 64.5 21 1 US-08-286-152-26 43 14.2 64.5 21 2 US-08-076-298-26 44 14.2 64.5 21 3 US-08-286-298-26 45 14.2 64.5 21 3 US-08-286-596-26 46 14.2 64.5 21 3 US-08-479-737-24 47 14.2 64.5 21 3 US-08-479-737-24 48 14.2 64.5 21 3 US-08-479-737-24
Length 22; ; Indels 0; Gaps 0;	25	Sequence 9, Appli Sequence 50, Appli Sequence 2, Appli Sequence 24, Appli Sequence 24, Appli Sequence 27, Appli Sequence 51, Appli Sequence 2, Appli Sequence 2, Appli Sequence 1, Appli Sequence 13, Appli Sequence 26, Appli

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TCGCACCCATCTCTCTCCTTCT 22

RESULT 2

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; Sequence 7, Application US/08178660
; Patent No. 5627277
; GENERAL INFORMATION:
APPLICANT: Cohen, Aharon S.
APPLICANT: Bourque, Andre
APPLICANT: Vilenchik, Maria
TITLE OF INVENTION: Method for Analyzing
TITLE OF INVENTION: Oligonucleotide Analogs
NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
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US-08-178-660-7
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NAME: KETHEY, AND-LOUISE
REGISTRATION NUMBER: 33,523
REFERENCE/DOCKET NUMBER: HYZ-
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/330-1300
TELEPAX: 617/330-1311
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UMBER: PCT/US95/00163
FILING DATE:
CLASSIFICATION:
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APPLICANT: Bourque, Andre
APPLICANT: Vilenchik, Maria
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ADDRESSEE: Lappin & Kusmer
STREET: 200 State Street
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                 STATE: N
COUNTRY:
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TYPE: nucleic acid
STRANDEDNESS: single
                                                         CITY: Boston
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02109
                                   Massachusetts
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                                                                        200 State Street
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RESULT 4
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                                                                                                                                                                                      COMPUTER: FLORDY disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PAtentin Release #1.0, Ver
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/00163
FILING DATE:
CLASSIFICATION:
ATTORNEY/AGFROR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 7, Application PC/TUS9500163
GENERAL INFORMATION:
APPLICANT: Cohen, Aharon S.
APPLICANT: Bourque, Andre
APPLICANT: Vilenchik, Maria
TITLE OF INVENTION: Method for Analyzing
TITLE OF INVENTION: Oligonucleotide Analogs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 100.
21; Conservative
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 base pairs
                                                                  TELEFAX: 617/330-1311
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Kerner, Ann-Louise
REGISTRATION NUMBER: 33,523
REFERENCE/DOCKET NUMBER: HYZ-
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/330-1300
TELEPAX: 617/330-1311
                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: KETNET, ANN-LOUISE
REGISTRATION NUMBER: 33,523
REFERENCE/DOCKET NUMBER: HY
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
               SEQUENCE CHARACTERISTICS:
LENGTH: 21 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Lappin & Kusmer
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HYPOTHETICAL:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/178,660 FILING DATE: CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Lappan ADDRESSEE: Lappan Compress: 200 State Street
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                                                                                       TELEPHONE: 617/330-1300
TELEFAX: 617/330-1311
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STRANDEDNESS:
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Pred. No.
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Sequence,
Patent INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Cohen, Aharon S.
APPLICANT: Cohen, Andre
APPLICANT: Bourque, Andre
APPLICANT: Vilenchik, Maria
TITLE OF INVENTION: Method for Analyzing
TITLE OF INVENTION: Oligonucleotide Analogs
NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
ADDRESSEB: Lappin & Kusmer
STREET: 200 State Street
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                                                   RESULT 6
PCT-US95-00163-8
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                                                                                                                                                                                                                                                                                              Sequence 8, Application PC/TUS9
GENERAL INFORMATION:
APPLICANT: Cohen, Aharon S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 8, April 5627277
                                                                                                                                                          Matches
                                                                                                                                                                   Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
                                                                                                                                                                                                                     TOPOLOGY: 1:
MOLECULE TYPE:
HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/178,660
FILING DATE:
CLASSIFICATION: 536
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: CDNA HYPOTHETICAL: NO
  APPLICANT:
                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                     LENGTH:
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                                       Application PC/TUS9500163
                                                                                                                    GCACCCATCTCTCTCCTTCT 22
                                                                                                      GCACCCATCTCTCTCCTTCT 20
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Bourque, Andre
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100.0%; Pr/
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                                                                                                                                                       Score 20; DB; Pred. No. 5.
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                                                                                                                                                                     DB 1;
5.7;
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US-08-420-672-5
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Patent No. 6372427
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                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
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MOLECULE TYPE:
HYPOTHETICAL:
                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: KERIMET, ANN-LOUISE
REGISTRATION UNMBER: 33,523
REFERENCE/DOCKET NUMBER: HY
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                             APPLICANT: Kandimalla, Ekambar R.
APPLICANT: Agrawal, Sudhir
TITLE OF INVENTION: COOPERATIVE OLIGONUCLEOTIDES
NUMBER OF SEQUENCES: 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Vilenchik, Maria
TITLE OF INVENTION: Method for Analyzing
TITLE OF INVENTION: Oligonucleotide Analogs
NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: Lappin & Kusmer
STREET: 200 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS
               ATTORNEY/AGENT INFORMATION:
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                                  FILING DATE:
CLASSIFICATION: 514
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                                                                                                                                                                                                                                               STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS:
                                                                 APPLICATION NUMBER: US/08/420,672
                                                                                                                                                                                COUNTRY: U
                                                                                                                                                                                                                               CITY: Boston
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TELEFAX: 617/330-1311
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nucleic acid
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Kerner, Ann-Louise
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7: U.S.A.
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200 State Street
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                                                                                                 Version
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Gaps

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REGISTRATION NUMBER:

33,523

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US-08-098-945A-1
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  Best Local Similarity
                      Query Match
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Best Local Similarity
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TELERAX: 617-330-1300
TELERAX: 617-330-1311
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                     TELEFAX: (617) 526-5000 INFORMATION FOR SEQ ID NO: 1: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       HYPOTHETICAL:
ANTI-SENSE: 1
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                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/08/098,945A
FILING DATE: 28-Jul-1993
CLASSIFICATION: CUNKnown>
ATTORNEY/AGENT INFORMATION:
NAME: Kerner, Ann-Louise
REGISTRATION NUMBER: 33,523
REFERENCE/DOCKET NUMBER: 33,523
REFERENCE/DOCKET NUMBER: HYS-498(47508-498)
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 21 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLIANAL APPRIANAL, Sudhir
Agrawal, Sudhir
TITLE OF INVENTION: Building Blocks with Carbamate
Internucleoside Linkages and No. 6605708el Oligonucleotides
Derived Therefrom
                                                                                                                   TOPOLOGY: linear MOLECULE TYPE: cDNA HYPOTHETICAL: NO
                                                                              SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                                                                                                     ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Habus, Ivan
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20;
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                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STATE: Massachusetts
COUNTRY: United States
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Boston
                                                                                                                                                                                                                                                                                                      TELEPHONE: (617) 526-6000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 02109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Hale and Dorr LLP
STREET: 60 State Street
                                                                                                                                                                                                                     LENGTH: 21 base pairs
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100.0%; Pred. No.
90.9%; Score 20; DB 4; 100.0%; Pred. No. 5.7;
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                    Length 21;
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US-09-167-375-24
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TITLE OF INVENTION: methods for their use.

FILE REFERENCE: MNGJ 2002 (CIP-1)

CURRENT APPLICATION NUMBER: US/09/167,375B

CURRENT FILING DATE: 1998-10-06

NUMBER OF SEQ ID NO 24

LENGTH: 22

TYPE: RNA

ORGANISM: Human Immunodeficiency Virus type 1

FEATURE:

FEATURE: Testing of the content of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 13; Conserv
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Patent No. 6207455
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                                                                                                                                                                                                                       COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Vei
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/935,312
FILING DATE: 22-SEP-1997
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: COOPER, Iver P.
REGISTRATION NUMBER: 28,005
REGISTRATION NUMBER: CHANG=112
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: CHANG, Lung-Ji
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT:
                                                                                                                             TELEFAX: 202-737-3528
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LOCATION: Portion of the gag gene OTHER INFORMATION: Complementary to a portion of the
                     SEQUENCE CHARACTERISTICS
LENGTH: 22 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
                                                                                                                                                                 TELEPHONE: 202-628-5197
TELEFAX: 202-737-3528
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 20001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET:
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE:
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Pred. No. 5.8;
7; Mismatches
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US-08-420-672-6
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Sequence 8, Application US/09318138
PATENT NO. 6531123
GENERAL INFORMATION:
APPLICANT: CHANG, Lung-Ji
TITLE OF INVENTION: LENTIVIRAL VECTORS
                                                                                                                                                                                          Matches
                                                                                                                                                                                                                      Query Match
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APPLICANT: Kandimalla, Ekambar R.
APPLICANT: Agrawal, Sudhir
TITLE OF INVENTION: COOPERATIVE OLIGONUCLEOTIDES
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-330-1300
TELEPAX: 617-330-1311
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
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MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = ""DNA""
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/420,672
FILING DATE:
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                             HYPOTHETICAL:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                              MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                         STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Kerner, Ann-Louise
REGISTRATION NUMBER: 33,523
REFERENCE/DOCKET NUMBER: HY
                                                                                                                                                                                         Local Similarity 95.2
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200 State Street
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95.2%;
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Pred. No. 11
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Pred. No. 1
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US-08-319-823A-10
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               ZIP: 60606

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: LEM PC compatible
OPERATING SYSTEM: Windows 95
SOFTWARE: Microsoft Word for Windows 95
CURRENT APPLICATION DATA:
ADDITICATION NUMBER: US/08/319,823A
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                                                                                                                                                                                                                                                                     APPLICANT: Agrawal, Sud
TITLE OF INVENTION: The:
TITLE OF INVENTION: Old:
TITLE OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
ADDRESCEPT.
                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: Agrawa
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                                                                                                                                                                                                        CITY: Chicago
STATE: Illinois
CLASSIFICATION:
                                                                                                                                                                                    COUNTRY:
                                                                                                                                                                                                                                             STREET:
                                                                                                                                                                                                                                                               ADDRESSEE:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/318,138
FILING DATE: 25-May-199
CLASSIFICATION: <UNKnown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US 60/086,635
PILING DATE: 26-MAY-1998
APPLICATION NUMBER: US 08/935,312
PILING DATE: 22-SEP-1997
ATTORNEY/AGENT INFORMATION:
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LENGTH: 22 base pairs
TYPE: nucleic acid
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REGISTRATION NUMBER: 28,005
REFERENCE/DOCKET NUMBER: CH
TELECOMMUNICATION INFORMATION:
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ADDRESSEE: BROWDY AND NEIMARK, P.L.L.C.
STREET: 624 Ninth Street, N.W., Suite 300
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MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                           300 South Wacker
                                                                                                                                                                                        USA
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                                                                                                                                                                                                                                         McDonnell Boehnen Hulbert & Berghoff, Ltd
10 South Wacker Drive, Seventh Floor
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                                                                                                                                                                                                                                                                                                                     Therapeutic Anti-HIV Oligonucleotides and
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Pred. No. 11;
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ATTORNEY/AGENT INFORMATION:

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Matches
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Patent No. 6489464
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Meschwitz, Susan M.
APPLICANT: Agrawal, Sudhir
TITLE OF INVENTION: Branched Oligonucleotides as Pathogen
TITLE OF INVENTION: Inhibitory Agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                    TELEFAX: (312)715-1239
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: Greenfield, Michael S.
REGISTRATION NUMBER: 37,142
REFERENCE/DOCKET NUMBER: 93,71
TELECOMMUNICATION INFORMATION:
TELEPHONE: (312)715-1000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: 312-913-0001
TELEFAX: 312-913-9808
INFORMATION FOR SEQ ID NO: 1
                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IN PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WordPerfect for DOS 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 19
CORRESPONDENCE ADDRESS:
ADDRESSEE: Michael S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/252,494
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE:
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LENGTH: 20 base pairs TYPE: nucleic acid STRANDEDNESS: single TOPOLOGY: linear
                                                                       APPLICATION NUMBER: FILING DATE: CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Chicago
                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET:
                                                                                                                                                                                                   FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LOCATION:
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chael S.
37,142
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US-08-664-786-2
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                                                        Matches
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                                                                       Best Local
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                                                                                                                                                                                                                                                                                  TELEFAX: (312)715-1234
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: Greenfield, Michael S.
REGISTRATION NUMBER: 37,142
REFERENCE/DOCKET NUMBER: 93,7
TELECOMMUNICATION INFORMATION:
TELEPHONE: (312)715-1000
                                                                                                                        FEATURE:

NAME/KEY: misc feature
LOCATION: 1.20
OTHER INFORMATION: /note
OTHER INFORMATION: inter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WordPerfect for DOS 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/664,786
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 60606
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 514
PRIOR APPLICATION NUMBER: 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
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                                                                                                                                                                                                   ANTI-SENSE:
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TELEPHONE: (312)715-1234
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE:
                                                                                                                                                                                                                                           STRANDEDNESS:
                                                                                                                                                                                                                                                         TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Chicago
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les 19; Conserv
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                          1 TCGCACCCATCTCTCTCT 19
                                                                       Similarity
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TCGCACCCATCTCTCTCCT 20
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10 S. Wacker Drive Suite 3000
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                                                        Conservative
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100.0%; Pred. No. 16
Live 0; Mismatches
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                                                                      86.4%; Score 19; 100.0%; Pred. No.
                                                                                                                             /note= "All phosphorothioate
internucleotide linkages"
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                                                       0;
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                                                                       DB 4; Length 20; 16;
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16;
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Search completed: September 14, 2005, 23:12:11

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Job time : 71.6047 secs

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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Scoring table:
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                   IDENTITY_NUC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Published Applications NA:*
1: /cgn2_6/ptodata/1/pubpna/US07
: /cgn2_6/ptodata//pubpna/PCT_NEW PUB.seq:*
:/cgn2_6/ptodata/I/pubpna/USO6_NEW PUB.seq:*
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:/cgn2_6/ptodata/I/pubpna/USO7_NEW PUB.seq:*
:/cgn2_6/ptodata/I/pubpna/USO7_NEW PUB.seq:*
:/cgn2_6/ptodata/I/pubpna/USO8_PUBCOMB.seq:*
:/cgn2_6/ptodata/I/pubpna/USO8_PUBCOMB.seq:*
:/cgn2_6/ptodata/I/pubpna/USO8_PUBCOMB.seq:*
:/cgn2_6/ptodata/I/pubpna/USO9A_PUBCOMB.seq:*
:/cgn2_6/ptodata/I/pubpna/USO9A_PUBCOMB.seq:*
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:/cgn2_6/ptodata/I/pubpna/USO0F_PUBCOMB.seq:*
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Copyright (c) 1993 - 2005 Compugen Ltd.
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500.694 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

7654827	Result No.
22 21 21 21 21	1
95.5 95.5 95.5 95.5 95.5	Query Match 1
21 21 22 22 22 21 21 21 21 21 21 21 21 2	Query Match Length DB ID
10 11 0 11 0 11 0 11 0 11 0 11 0 11 0	B 11
US-09-896-692B-5 US-09-896-692B-1 US-09-896-692B-2 US-09-896-692B-3 US-09-896-692B-3 US-09-896-692B-4 US-09-896-692B-6	query Match Length DB ID
Sequence 5, Appli Sequence 35, Appli Sequence 1, Appli Sequence 2, Appli Sequence 3, Appli Sequence 4, Appli Sequence 6, Appli	Description

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-10-157-580A- -10-157-580A- -10-157-580A- -10-157-580A-	US-10-923-473-409 US-10-923-473-463 US-10-923-473-1147 US-10-923-473-1201 US-10-157-580A-26 US-10-157-580A-41	-10-640-898 -10-054-4298 -10-483-654 -10-225-023 -10-225-023 -10-225-023	-10-225 -10-225 -10-225 -10-923 -10-923 -10-923 -10-923 -10-923 -10-923 -10-923	-10-262-318-1 -10-054-429A -10-054-429A -10-225-023-4 -10-225-023-4 -10-225-023-4 -10-225-023-4
Sequence 42, Appl Sequence 61, Appl Sequence 69, Appl Sequence 73, Appl		e 463,		e 11, App e 5, App e 6, App e 420, App e 429, App e 440, App e 440

ALIGNMENTS

RESULT 1 US-09-896-692B-5

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                                                                                                                                                 SEQ ID NO 5
LENGTH: 22
TYPB: DNA
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: synthetic modified antisense oligonucleotide
                                                                                                                                                                                                                                                                        Sequence 5, Application US/09896692B
Publication No. US20030100521A1
GENERAL INFORMATION:
APPLICANT: Agrawal, Sudhir
TITLE OF INVENTION: No. US20030100521A1e1 HIV-Specific Synthetic Oligonucleotides and
TITLE OF INVENTION: Methods of Their Use
FILE REFERENCE: 47508.556 (HYZ-069)
CURRENT APPLICATION NUMBER: US/09/896,692B
CURRENT FILING DATE: 2002-10-17
PRIOR FILING DATE: 1997-08-19
NUMBER OF SEQ ID NOS: 8
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                       Matches
                                                                                         Best Local Similarity
                                                                                                          Query Match
22; Conservative
                                                                                       100.0%; Score 22; DB 10; 100.0%; Pred. No. 4.3;
                                                                     0; Mismatches
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PRIOR FILING DATE: 2001-01-19
PRIOR APPLICATION NUMBER: US 09/487,023
PRIOR FILING DATE: 2000-01-19
PRIOR APPLICATION NUMBER: US 09/016,541
PRIOR FILING DATE: 2000-11-24
PRIOR FILING DATE: 1998-01-30
PRIOR FILING DATE: 1998-01-30
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                                               Matches
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 35
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                             APPLICANT: Agrawal, Sudhir
TITLE OF INVENTION: No. US20030100521A1el HIV-Specific Synthetic Oligonucleotides
TITLE OF INVENTION: Methods of Their Use
FILE REFERENCE: 47508.556 (HYZ-069)
CURRENT APPLICATION NUMBER: US/09/896,692B
CURRENT FILING DATE: 2002-10-17
                                                                                                                                                                                                                                                                                 PRIOR APPLICATION NUMBER: US 08/914,827
PRIOR FILING DATE: 1997-08-19
NUMBER OF SEQ ID NOS: 8
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APPLICANT: Masood, Rizwan
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR .
TITLE OF INVENTION: VEGF OLIGONUCLEOTIDES
FILE REFERENCE: 21327-701CON2
CURRENT APPLICATION NUMBER: US/09/805,761
CURRENT FILING DATE: 2001-03-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: US 60/037,004 PRIOR FILING DATE: 1997-01-31
                                                                                                                                        OTHER INFORMATION: synthetic modified antisense oligonucleotide
                                                                                                                                                                              ORGANISM: Artificial Sequence
                                                                                                                                                                                                LENGTH: 21
TYPE: DNA/RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: nucleotides 1-4 and 1 OTHER INFORMATION: methylribonucleosides
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                                                                                                                                                                 FEATURE:
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                                                                Local
                                           l Similarity
18; Conserv
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    TCGCACCCATCTCTCTCCTTC 21
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                                             Conservative
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                                                            95.5%;
85.7%;
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Pred. No.
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Pred. No.
                                             Mismatches
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                                                            DB
12;
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                                                                                Length 21
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FILE REFERENCE: 47508.556 (HYZ-069)
CURRENT APPLICATION NUMBER: US/99/896,692B
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: US 08/914,827
PRIOR FILING DATE: 1997-08-19
NUMBER OF SEQ ID NOS: 8
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 3
SEQ ID NO 3
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                                                            RESULT 6
US-09-896-692B-4
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US-09-896-692B-3
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US-09-896-692B-2
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Sequence 4, Application US/09896692B Publication No. US20030100521A1 GENERAL INFORMATION:
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Publication No. US20030100521A1
GENERAL INFORMATION:
APPLICANT: Agrawal, Sudhir
TITLE OF INVENTION: No. US20030100521A1el HIV-Specific Synthetic Oligonucleotides and
TITLE OF INVENTION: Methods of Their Use
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 2
LENGTH: 21
TYPE: DNA
ORGANISM: Artificial Sequence
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APPLICANT: Agrawal, Sudhir
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CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: US 08/914,827
PRIOR FILING DATE: 1997-08-19
NUMBER OF SEQ ID NOS: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Agrawal, Sudhir
TITLE OF INVENTION: No. US20030100521A1el HIV-Specific Synthetic Oligonucleotides and
TITLE OF INVENTION: Methods of Their Use
FILE REFERENCE: 47508.556 (HYZ-069)
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                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: synthetic modified antisense oligonucleotide
                                                                                                                                                                                                                                                                                                                                                ORGANISM: Artificial Sequence FEATURE:
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                                                                                                                                                                                                                                             Local Similarity
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85.7%;
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Pred. No.
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Pred. No. 12;
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RESULT 8
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APPLICANT: Copharos
APPLICANT: Collins, Douglas A.
TITLE OF INVENTION: CODMINISTRATION OF TRANS
TITLE OF INVENTION: DELIVER AGENTS
FILE REFERENCE: COP1012
CURRENT APPLICATION NUMBER: US/10/262,318
CURRENT FILING DATE: 2002-09-30
NUMBER OF SEQ ID NOS: 14
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; OTHER INFORMATION: synthetic modified oligonucleotide US-09-896-692B-6
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; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: synthetic modified antisense oligonucleotide
US-09-896-692B-4
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TITLE OF INVENTION: No. US20030100521A1el HIV-Specific Synthetic Oligonucleotides
TITLE OF INVENTION: Methods of Their Use
FILE REFERENCE: 47508.556 (HYZ-069)
CURRENT FILIG DATE: 2002-10-17
PRIOR FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: US 08/914,827
PRIOR APPLICATION NUMBER: US 08/914,827
PRIOR FILING DATE: 1997-08-19
NUMBER OF SEQ ID NOS: 8
SOFTWARE: FastSEQ for Windows Version 4.0
LENCTH. 21
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Best Local
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Best Local 9
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TITLE OF INVENTION: No. US20030100521A1el HIV-Specific Synthetic Oligonucleotides and
TITLE OF INVENTION: Methods of Their Use
FILE REFERENCE: 47508.556 (HYZ-069)
CURRENT APPLICATION NUMBER: US/09/896,692B
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: US 08/914,827
PRIOR FILING DATE: 1997-08-19
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SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
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                                                                                                                                           TRANSPORT PROTEINS WITH CONJUGATED COBALAMIN
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APPLICANT: Agrawal, Sudhir
ITILE OF INVENTION: Cooperative Oligonucleotides
FILE REFERENCE: 47508-580 (HYZ-027CIP)
CURRENT APPLICATION NUMBER: US/10/054,429A
CURRENT FILING DATE: 2002-11-04
PRIOR APPLICATION NUMBER: US 08/420,672
PRIOR FILING DATE: 1995-04-12
NUMBER OF SEQ ID NOS: 30
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 5
LENGTH: 21
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                                                                                                        GENERAL INFORMATION:
                                                                                                                          Sequence 6, Application US/10054429A Publication No. US20030099959A1
                                                                                                                                                                                                                                                                                                      Matches
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Best Local Similarity
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LENGTH: 21
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Publication No. US20030099959A1
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APPLICANT: Agrawal, Sudhir
TITLE OF INVENTION: Cooperative Oligonucleotides
FILE REFERENCE: 47508-580 (HYZ-027CIP)
CURRENT APPLICATION NUMBER: US/10/054,429A
CURRENT FILING DATE: 2002-11-04
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                                                                          APPLICANT: Kandimalla, Ekambar R. APPLICANT: Agrawal, Sudhir
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                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
ORGANISM: Artificial Sequence
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OTHER INFORMATION: 2'OMe
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OTHER INFORMATION: Oligonucleotide-- GEM-92 (Hybridon)
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85.7%;
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Pred. No.
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PRIOR APPLICATION NUMBER: US 08/420,672
PRIOR FILING DATE: 1995-04-12
NUMBER OF SEQ ID NOS: 30
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 6

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APPLICANT: McSwiggen, James
TITLE OF INVENTION: RNA Interference Mediated Inhibition of
TITLE OF INVENTION: Interfering RNA
TITLE OF INVENTION: Interfering RNA
FILE REFERENCE: 400/054 (MEHBO1-665-B)
CURRENT APPLICATION NUMBER: US/10/225,023
CURRENT FILING DATE: 2003-01-06
PRIOR APPLICATION NUMBER: US 60/398,036
PRIOR APPLICATION NUMBER: US 60/398,036
PRIOR FILING DATE: 2002-07-23
PRIOR FILING DATE: 2002-05-29
PRIOR APPLICATION NUMBER: US 60/294,140
PRIOR FILING DATE: 2002-05-29
PRIOR APPLICATION NUMBER: US 10/157,580
PRIOR APPLICATION NUMBER: US 10/157,580
PRIOR FILING DATE: 2002-05-29
NUMBER OF SEQ ID NOS: 1494
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; OTHER INFORMATION:
US-10-054-429A-6
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Publication No. US20030175950A1
GENERAL INFORMATION:
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Best Local Similarity
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Best Local Similarity
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CURRENT FILING DATE: 2003-01-06
PRIOR APPLICATION NUMBER: US 60/398,036
PRIOR FILING DATE: 2002-07-23
PRIOR APPLICATION NUMBER: US 60/294,140
PRIOR FILING DATE: 2002-05-29
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TITLE OF INVENTION: RNA Interference Mediated Inhibition of HIV Gene Expression Using
TITLE OF INVENTION: Interfering RNA
FILE REFERENCE: 400/054 (MBHB01-665-B)
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SOFTWARE: PatentIn version 3.0
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ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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o. US20030175950A1
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Pred. No. 59;
0; Mismatches
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RESULT 14
US-10-225-023-466/c
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US-10-225-023-440/c
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Matches
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LENGTH: 19
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                                                  TITLE OF INVENTION: RNA Interference Mediated Inhibition of HIV Gene Expression Using TITLE OF INVENTION: Interfering RNA FILE OF INVENTION: Interfering RNA FILE REFERENCE: 400/054 (MBHB01-665-B) CURRENT APPLICATION NUMBER: US/10/225,023 CURRENT FILING DATE: 2003-01-06 PRIOR APPLICATION NUMBER: US 60/398,036 PRIOR PELICATION NUMBER: US 60/398,036 PRIOR PILLING DATE: 2002-07-23 PRIOR PILLING DATE: 2002-07-23 PRIOR PILLING DATE: 2002-05-29 PRIOR FILING DATE: 2002-05-29
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CURRENT FILING DATE: 2003-01-06
PRIOR APPLICATION NUMBER: US 60/398,036
PRIOR FILING DATE: 2002-07-23
PRIOR APPLICATION NUMBER: US 60/294,140
PRIOR FILING DATE: 2002-05-29
PRIOR FILING DATE: 2002-05-29
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ORGANISM: Artificial Sequence
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TYPE: RNA
ORGANISM: Artificial Sequence
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              APPLICATION NUMBER: US 10/157,580 FILING DATE: 2002-05-29
OF SEQ ID NOS: 1494
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No. US20030175950A1
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APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: McSwiggen, James
TITLE OF INVENTION: Interference Mediated Inhibition of HIV Gene Expression Using
TITLE OF INVENTION: Interfering RNA
FILE REFERENCE: 400/054 (MBHB01-665-B)
CURRENT FILING NOWEBR: US/10/225,023
CURRENT FILING DATE: 2003-01-06
PRIOR APPLICATION NUMBER: US 60/398,036
PRIOR FILING DATE: 2002-07-23
PRIOR FILING DATE: 2002-05-29
PRIOR FILING DATE: 2002-05-29
PRIOR APPLICATION NUMBER: US 10/157,580
PRIOR APPLICATION SEQ ID NOS: 11494
SOFTWARE: Patentin version 3.0
SEQ ID NO 1158
LEEGTH: 19
TYPE: RNA
ORGANISM: Artificial Sequence
PEATURE:
OTHER INFORMATION: Description of Artificial Sequence: siNA antisense region
US-10-225-023-1158
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; SEQ ID NO 466
; SEQ ID NO 465
; LENCTH: 19
; TYPE: RNA
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Target sequence/siNA sense
US-10-225-023-466
Search completed: September 15, 2005, 00:50:14 Job time: 288.558 secs
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US-10-225-023-1158
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Best Local Similarity
Matches 13; Conserv
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Post-processing: Minimum Match 0%
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m2_6/ptodata/1/pna/US080_COMB.seq:*
m2_6/ptodata/1/pna/US080_COMB.seq:*
m2_6/ptodata/1/pna/US080_COMB.seq:*
m2_6/ptodata/1/pna/US082_COMB.seq:*
m2_6/ptodata/1/pna/US083_COMB.seq:*
m2_6/ptodata/1/pna/US083_COMB.seq:*
m2_6/ptodata/1/pna/US083_COMB.seq:*
m2_6/ptodata/1/pna/US085_COMB.seq:*
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m2_6/ptodata/1/pna/US085_COMB.seq:*
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26/ptodata/1/pna/PCTUS1 COMB.seq:
26/ptodata/1/pna/PCTUS2 COMB.seq:
26/ptodata/1/pna/PCTUS COMB.seq:*
26/ptodata/1/pna/US06 COMB.seq:*
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15: /cgm2_6/prodata/1/pna/US099F_COMB.seq;*
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19: /cgm2_6/prodata/1/pna/US101C_COMB.seq;*
10: /cgm2_6/prodata/1/
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Result No.

Score

Query Match Length

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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/cgm2_6/ptodata/1/pna/US6058_COMB.seq:*
/cgm2_6/ptodata/1/pna/US6059_COMB.seq:*
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5. US-08-914-827-1
5. US-08-914-827-2
5. US-08-914-827-3
5. US-08-914-827-4
5. US-08-914-827-6
6. US-09-837-806-1
6. US-09-837-806-2
6. US-09-837-806-2
6. US-09-837-806-3
6. US-09-837-806-3
6. US-09-837-806A-1
6. US-09-837-806A-1
6. US-09-837-806A-1
7. US-09-837-806A-1
8. US-09-837-806A-2
8. US-09-837-806A-3
8. US-09-837-806A-3
8. US-09-896-692B-3
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                                                       ALIGNMENTS
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                                                                                                                                                                                                                                                              Sequence 5, Application US/08914827
GENERAL INFORMATION:
APPLICANT: Agrawal, Sudhir
TITLE OF INVENTION: NOVEL HIV-SPE
TITLE OF INVENTION: OLIGONUCLEOTI
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSEE: HALE AND DORR LLP
STREET: 60 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-08-478-464-6
US-08-478-464-6
Sequence 6, Application US/08478464
; GENERAL INFORMATION:
APPLICANT: Cohen, Aharon S.
APPLICANT: Bourque, Andre
APPLICANT: Vilenchik, Maria
TITLE OF INVENTION: Method for Analyzing
TITLE OF INVENTION: Oligonucleotide Analogs
NUMBER OF SEQUENCES: 8
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SEQUENCE CHARACTERISTICS:
LENGTH: 22 base pairs
TYPE: nucleic acid
TYPE: single
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US-08-914-827-5
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Best Local S
Matches 22
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
COMPUTER: PATENTIN PC-DOS/MS-OOS
SOFTWARE: PATENTIN Release #1.0, Vers
SOFTWARE: PATENTIN NUMBER: US/08/478,464
FILING DATE: 07-UNB-1995
CLASSIFICATION: 536
ATTORNEY/AGENT INFORMATION:
NAME: Kerner, Ann-Louise
REGISTRATION NUMBER: 33,523
REFERENCE/DOCKET NUMBER: HYZ-012DV
                                                                                                      COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
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TELEPHONE: 617/330-1300
TELEPAX: 617/330-1311
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ADDRESSEE: Lappin & Kusmer
STREET: 200 State Street
CURRENT APPLICATION DATA:
                    COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: 200 State St:
CITY: Boston
STATE: Massachusetts
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CITY: Boston
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ATTORNEY/AGENT INFORMATION:

CLASSIFICATION:

APPLICATION NUMBER: FILING DATE:

US/08/914,827

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GENERAL INFORMATION:
GENERAL THEORMATION: NOVEL HIV-SPECIFIC SYNTHETIC
OLIGONUCLEOTIDES AND METHODS OF THEIR USE
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Matches
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                                                                                                                           INFORMATION FOR SEQ ID NO: 5
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REGISTRATION NUMBER: 33,523
REFERENCE/DOCKET NUMBER: HY
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 526-6000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ANTI-SENSE: YES
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                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: 009/837,806
FILING DATE: 30-Mar-2001
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                        APPLICATION NUMBER: 08/914,827
FILING DATE: «Unknown»
ATTORNEY/AGENT INFORMATION:
NAME: Kerner, Ann-Louise
REGISTRATION NUMBER: 33,523
REFERENCE/DOCKET NUMBER: HYZ-069
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 526-6000
TELEPAX: (617) 526-5000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS: 811
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TYPE: nucleic acid
STRANDENNESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ENGTH:
                                                                                                        SEQUENCE CHARACTERISTICS:
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                                                                                       LENGTH: 22 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: United States Of America
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100.0%; Pred. No.
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; OTHER INFORMATION: Description of Artificial Sequence: Synthetic
; OTHER INFORMATION: oligonucleotide
US-09-837-806A-5
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US-09-837-806A-5
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                                                                                                                                     ; ORGANISM: Artificial Sequence; FEATURE:
; PEATURE:
; OTHER INFORMATION: synthetic modified antisense oligonucleotide
US-09-896-692B-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 5
US-09-896-692B-5
                                                                                                                                                                                                      SEQ ID NO 5
LENGTH: 22
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOPTWARE: PatentIn Ver. 3.2
SEQ ID NO 5
SENGTH: 22
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 5, Application US/09896692B GENERAL INFORMATION:
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Best Local Similarity
Matches 22; Conserv
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                                                                                  Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                   APPLICANT: Agrawal, Sudhir
TITLE OF INVENTION: Novel HIV-Specific Synthetic Oligonucleotides and
TITLE OF INVENTION: Methods of Their Use
FILE REFERENCE: 47508.556 (HYZ-069)
CURRENT APPLICATION NUMBER: US/09/96,692B
CURRENT APPLICATION NUMBER: US/09/96,692B
CURRENT FILING DATE: 2002-10-17
PRIOR APPLICATION NUMBER: US 08/914,827
PRIOR PILING DATE: 1997-08-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity
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PRIOR FILING DATE: 1997-08-19
NUMBER OF SEQ ID NOS: 7
SOFTWARE: Paranta
                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 8
SOFTWARE: FastSEQ for Windows Version 4.0
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CURRENT APPLICATION NUMBER: US/09/837,806A
CURRENT FILING DATE: 2001-04-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: NOVEL HIV-SPECIFIC SYNTHETIC OLIGONUCLEOTIDES AND TITLE OF INVENTION: METHODS OF THEIR USE
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                                TCGCACCCATCTCTCTCTTCT 22
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100.0%; Pred. No. 32;
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Pred. No.
                                                                                   Score 22; DE
Pred. No. 32;
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RESULT 7
US-08-914-827-1
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GENERAL INFORMATION:
APPLICANT: Cohen, Aharon S.
APPLICANT: Bourque, Andre
APPLICANT: Vilenchik, Maria
TITLE OF INVENTION: Method for Analyzing
TITLE OF INVENTION: Oligonucleotide Analogs
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TELEFAX: 617/330-1311
INFORMATION FOR SEQ ID NO:
ZIP: 02109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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MEDIUM TYPE: FORDy disk
COMPUTER: IBM PC compatible
COMPUTER: PC-DOS/MS-DOS
SOFTMARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/478,464
FILING DATE: 07-UUNE-1995
CLASSIFICATION: 536
CLASSIFICATION: 536
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NAME: KETHEY, AND-LOUISE
REGISTRATION NUMBER: 33,523
REFERENCE/DOCKET NUMBER: HY
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                     NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
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TITLE OF INVENTION: NOVEL HIV-SPECIFIC SYNTHETIC
TITLE OF INVENTION: OLIGONUCLEOTIDES AND METHODS OF THEIR USE
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LENGTH: 21 base pairs
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CITY: Boston
STATE: MA
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COUNTRY:
                                                                            COUNTRY: UZIP: 02109
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100.0%;
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US-08-914-827-2
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GENERAL INFORMATION:
APPLICANT: Agrawa
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Best Local Similarity
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NAME: KETNET, Ann-Louise
REGISTRATION UNMER: 33,523
REFERENCE/DOCKET NUMBER: HYZ-069
TELECOMUNICATION INFORMATION:
TELEPROUS: (617) 526-6000
TELEPRAX: (617) 526-5000
                                                                                                                                                                                 CLASSIFICATION: 514
ATTORNEY AGENT INFORMATION:
NAME: KEYDET, AND-LOUISE
REGISTRATION NUMBER: 33,523
REFERENCE/DOCKET NUMBER: HYZ-
TELECOMMUNICATION INFORMATION:
TELECHIONE: (617) 526.6000
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              MOLECULE TYPE:
HYPOTHETICAL: 1
ANTI-SENSE: YE
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MEDIUM TYPE: Floppy disk
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HYPOTHETICAL: N
ANTI-SENSE: YES
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ADDRESSEE: HALE AND DORR LLP
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LENGTH: 21 base pairs
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                                                                               LENGTH: 21 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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OPERATING SYSTEM: PC-DOS/MS-DOS
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85.7%;
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                                                         Sequence 4, Application US/08914827
GENERAL INFORMATION:
APPLICANT: Agrawal, Sudhir
TITLE OF INVENTION: NOVEL HIV-SPECIFIC SYNTHETIC
TITLE OF INVENTION: OLIGONUCLEOTIDES AND METHODS OF THEIR USE
NUMBER OF SEQUENCES: 6
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Best Local Similarity
Matches 21; Conserv
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TELEPHONE: (617) 526-6000
TELEPAX: (617) 526-5000
VEORMATION FOR SEQ ID NO: 3:
                                                                                                                                                                                                                                                                                                                                                ANTI-SENSE:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version
CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy
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TITLE OF INVENTION: NOVEL HIV-SPECIFIC SYNTHETIC
TITLE OF INVENTION: OLIGONUCLEOTIDES AND METHODS OF THEIR USE
NUMBER OF SEQUENCES: 6
                                            CORRESPONDENCE ADDRESS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Kerner, Ann-Louise
REGISTRATION NUMBER: 33,
REFERENCE/DOCKET NUMBER:
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CLASSIFICATION: 514
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              ADDRESSEE: HALE AND DOR'STREET: 60 State Street
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                          HALE AND DORR LLP
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Pred. No.
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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (617) 526-5000 INFORMATION FOR SEQ ID NO: 4
                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                      CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: HALE AND DORR LLP
                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Agrawal, TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                               TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 526-6000
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                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                             NAME: Kerner, Ann-Louise
REGISTRATION NUMBER: 33,
REFERENCE/DOCKET NUMBER:
                                                                                                                                     APPLICATION NUMBER: FILING DATE:
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REGISTRATION NUMBER: 33,523
REFERENCE/DOCKET NUMBER: HY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
CLASSIFICATION: 514
                TELEFAX:
                                                                                                                                                                                                                                                                     COUNTRY: U
ZIP: 02109
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                                                                                                                                                                                                                                                                                                                                STREET:
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TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/08/914,827
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                (617) 526-5000
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NOVEL HIV-SPECIFIC SYNTHETIC
OLIGONUCLECTIDES AND METHODS OF THEIR USE
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RESULT 12
US-09-708-786-2
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                                                                                                                                           PRIOR APPLICATION NUMBER: PCT/US01/00019
PRIOR FILING DATE: 2001-01-19
PRIOR PPLICATION NUMBER: US 09/487,023
PRIOR FILING DATE: 2000-01-19
PRIOR APPLICATION NUMBER: US 09/016,541
PRIOR PRICING DATE: 2000-11-24
PRIOR PRICING DATE: 1998-01-30
PRIOR FILING DATE: 1998-01-30
                                              SOFTWALL SEQ ID NO 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn version 3.0 SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 35, Application US/09805761 GENERAL INFORMATION:
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Best Local :
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TITLE OF INVENTION: WETHODS AND COMPOSITIONS FOR ANTISENSE
TITLE OF INVENTION: VEGF OLIGONUCLECTIDES
FILE REFERENCE: 21327-701CON2
CURRENT APPLICATION NUMBER: US/09/805,761
CURRENT FILING DATE: 2001-03-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Agrawal, Sudhir
TITLE OF INVENTION: Potentiation of Prodrug Efficacy
FILE REPERENCE: 47508.700US
CURRENT APPLICATION NUMBER: US/09/708,786
CURRENT FILING DATE: 2000-11-08
PRIOR APPLICATION NUMBER: PCT/US00/30687
PRIOR PILING DATE: 2000-11-08
NUMBER OF SEQ ID NOS: 3
                                                                               PRIOR FILING DATE: 1997-01-31
NUMBER OF SEQ ID NOS: 64
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                 PRIOR APPLICATION NUMBER: US 60/037,004
                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Gill, Parkesh APPLICANT: Masood, Rizw
TYPE: DNA
TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
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MOLECULE TYPE:
HYPOTHETICAL: N
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US-09-837-806-1
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GENERAL INFORMATION:
APPLICANT: Agrawal, Sudhir
TITLE OF INVENTION: NOVEL HIV-SPECIFIC SYNTHETIC
OLIGONUCLEOTIDES AND METHODS
                                                                   Matches
                                                                                                  Query Match
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Best Local Similarity
                                                                                  Best Local Similarity
                                                                                                                                                                                                                                                                                                        TELEPHONE: (617) 526-6000
TELEPAX: (617) 526-5000
INFORMATION FOR SEQ ID NO: 1:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ve:
CURRENT APPLICATION NUMBER: US/09/837,806
FILING DATE: 30-Mar-2001
                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 08/914,827
FILING DATE: «Unknown»
ATTORNEY/AGENT INFORMATION:
NAME: Kerner, Ann-Louise
REGISTRATION NUMBER: 33,523
REFERENCE/DOCKET NUMBER: HYZ-069
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                        SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                                                                                                                                                                         ANTI-SENSE: YES
                                                                                                                                                                                                        TOPOLOGY: linear MOLECULE TYPE: DNA/RNA
                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
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                 1 TCGCACCCATCTCTCTCTTC 21
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                                                                                                                                                                                                                                    LENGTH: 21 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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STREET: 60 State Street
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                                                                                  95.5%;
85.7%;
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Pred. No.
                                                                                  Score 21; DB 36;
Pred. No. 87;
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87;
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RESULT 15 US-09-837-806-2

Sequence 2, Application US/09837806 GENERAL INFORMATION:

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CITY: Boston
STATE: MA
COUNTRY: United States Of America
ZIP: 02109
COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: EMP CC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRINT APPLICATION NUMBER: US/09/837,806
FILING DATE: 30-Mar-2001
CLASSIFICATION NUMBER: US/09/837,806
FILING DATE: 4Unknown>
PRIOR APPLICATION NUMBER: 08/914,827
FILING DATE: CUNknown>
ATTORNEY/AGENT INPORMATION:
NAME: Kerner, Ann-Louise
REGISTRATION NUMBER: 33,523
REFERENCE/DOCKET N
Search completed: September 15, 2005, 00:40:44
Job time: 1770.21 secs
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                                                                                                                                                                                                                                                                                                                                           Query Match 95.5%; Score 21; DB 36; Length 21; Best Local Similarity 100.0%; Pred. No. 87; Matches 21; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: HALE AND DORR LLP
STREET: 60 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Agrawal, Sudhir
TITLE OF INVENTION: NOVEL HIV-SPECIFIC SYNTHETIC
OLIGONUCLECTIDES AND METHODS OF THEIR USE
NUMBER OF SEQUENCES: 6
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